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Chapter 1

dataField: totalPucksRatio (totalPucksRatio)

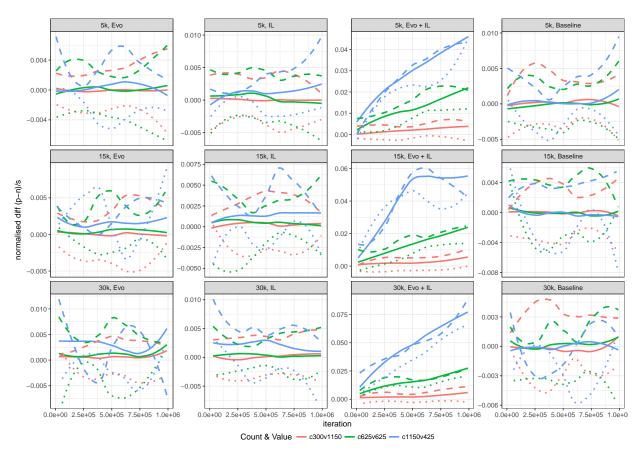
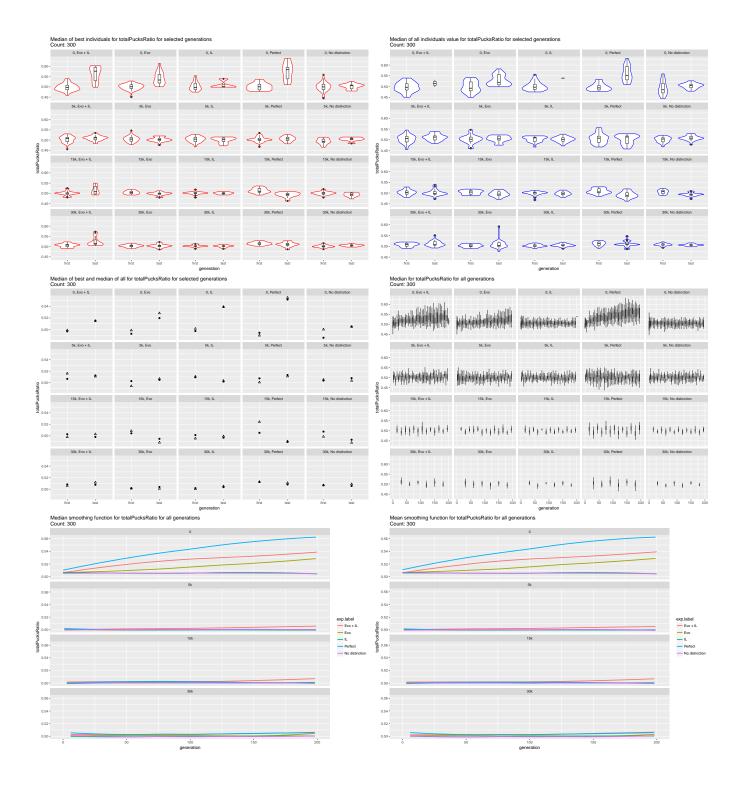


Figure 1.1: Normalised difference between positive and negative tokens collected. Solid line is value combined over all seasons, dashed = season 0, dotted = season 1

Chapter 2

Count: 300 Value: 1150



2.1 Experiments by Seasons. Count: 300 Value: 1150

2.1.1 Season: 0, Count: 300 Value: 1150

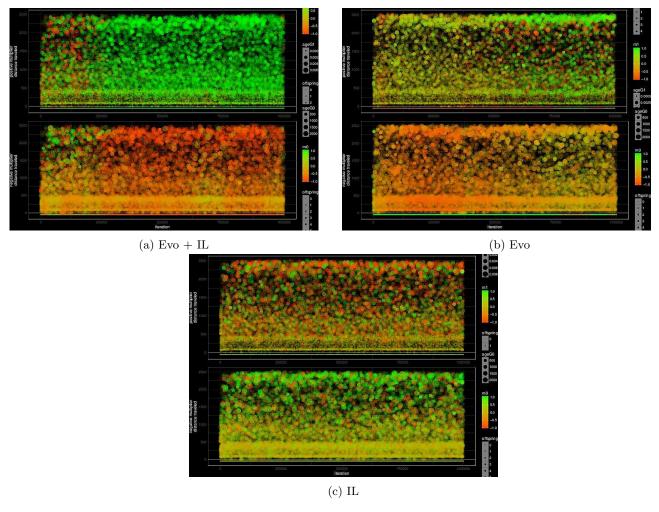


Figure 2.1: The graphs show multiple performance values to compare different adaptation mechanism: evolution combined with individual learning (\mathbf{a}), just evolution (\mathbf{b}) and just individual learning (\mathbf{c}). Data plotted shows 10% of the population (randomly sampled) and is taken from a single randomly selected run of the experiment. Each sub-graph shows the value of a single multiplier (names "positive" and "negative" refer to the initial value of the corresponding token), hence each genome has a data point on each of the sub-graphs. The values end-of-lifetime (x position), reproductive success (transparency; measured in number of offspring), adaptation success (size; measured in negative/positive token avoided/collected per life-time) and distance travelled (y-value) are the same in each sub-graph. The red/green lines below the y-axis is made up of dots, each representing the learning sign LS of the individuals: red=negative and green=positive. Note that LS is negated in the lower sub-graph.

	Evo	IL	Evo + IL	Baseline	Perfect
1	0.5306	0.5301	0.5034	0.5624	0.5029

Table 2.1: Median value of totalPucksRatio over the last 5000 iteration (N:30). Seasonal change every 0 iterations for configuration count=300 and value=1150.

	IL	Evo + IL	Baseline	Perfect
Evo	$> 6.04 \mathrm{e} ext{-}35$	= 6.64e-01	> 3.96e- 64	< 2.18e-32
IL		< 3.6e-26	= 1.64e-01	< 4.39 e-66
Evo + IL			> 1.02e-43	< 1.01e-18
Baseline				< 2.34 e-111

Table 2.2: Showing p-values of pairwise comparison of experiments for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Seasonal change every 0 iterations for configuration count=300 and value=1150. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

2.1.2 Season: 5k, Count: 300 Value: 1150

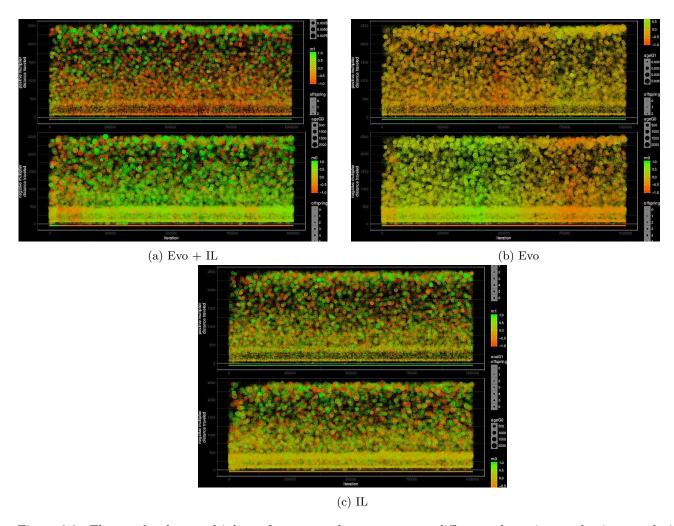


Figure 2.2: The graphs show multiple performance values to compare different adaptation mechanism: evolution combined with individual learning (\mathbf{a}), just evolution (\mathbf{b}) and just individual learning (\mathbf{c}). Data plotted shows 10% of the population (randomly sampled) and is taken from a single randomly selected run of the experiment. Each sub-graph shows the value of a single multiplier (names "positive" and "negative" refer to the initial value of the corresponding token), hence each genome has a data point on each of the sub-graphs. The values end-of-lifetime (x position), reproductive success (transparency; measured in number of offspring), adaptation success (size; measured in negative/positive token avoided/collected per life-time) and distance travelled (y-value) are the same in each sub-graph. The red/green lines below the y-axis is made up of dots, each representing the learning sign LS of the individuals: red=negative and green=positive. Note that LS is negated in the lower sub-graph.

	Evo	IL	Evo + IL	Baseline	Perfect
1	0.5029	0.4995	0.5006	0.5043	0.5013

Table 2.3: Median value of totalPucksRatio over the last 5000 iteration (N:30). Seasonal change every 5k iterations for configuration count=300 and value=1150.

	IL	Evo + IL	Baseline	Perfect
Evo	= 8.45e-01	$< 3.18 \mathrm{e} ext{-}05$	= 8.11e-02	= 5.38e-02
IL		< 1.27 e-04	= 1.17e-01	< 2.23 e-02
Evo + IL			> 2.18e-02	= 1.59e-01
Baseline				= 5.47e-01

Table 2.4: Showing p-values of pairwise comparison of experiments for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Seasonal change every 5k iterations for configuration count=300 and value=1150. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

2.1.3 Season: 15k, Count: 300 Value: 1150

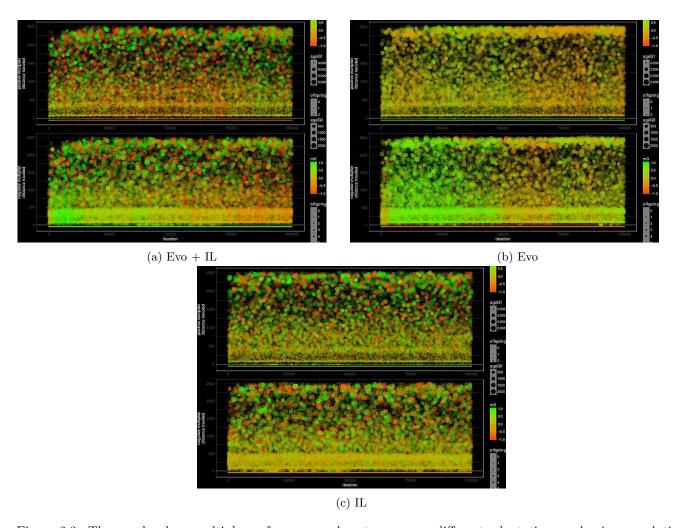


Figure 2.3: The graphs show multiple performance values to compare different adaptation mechanism: evolution combined with individual learning (\mathbf{a}), just evolution (\mathbf{b}) and just individual learning (\mathbf{c}). Data plotted shows 10% of the population (randomly sampled) and is taken from a single randomly selected run of the experiment. Each sub-graph shows the value of a single multiplier (names "positive" and "negative" refer to the initial value of the corresponding token), hence each genome has a data point on each of the sub-graphs. The values end-of-lifetime (x position), reproductive success (transparency; measured in number of offspring), adaptation success (size; measured in negative/positive token avoided/collected per life-time) and distance travelled (y-value) are the same in each sub-graph. The red/green lines below the y-axis is made up of dots, each representing the learning sign LS of the individuals: red=negative and green=positive. Note that LS is negated in the lower sub-graph.

	Evo	IL	Evo + IL	Baseline	Perfect
1	0.4981	0.4960	0.4973	0.4877	0.4929

Table 2.5: Median value of totalPucksRatio over the last 5000 iteration (N:30). Seasonal change every 15k iterations for configuration count=300 and value=1150.

	IL	Evo + IL	Baseline	Perfect
Evo	= 7.84e-02	$< 8.55 \mathrm{e} ext{-}04$	> 7.84e-04	> 9.88e-06
IL		= 5.27e-02	> 9.3e-09	> 5.65e-09
Evo + IL			> 1.37 e-14	$> 3.98 \mathrm{e} ext{-}13$
Baseline				= 5.94e-02

Table 2.6: Showing p-values of pairwise comparison of experiments for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Seasonal change every 15k iterations for configuration count=300 and value=1150. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

2.1.4 Season: 30k, Count: 300 Value: 1150

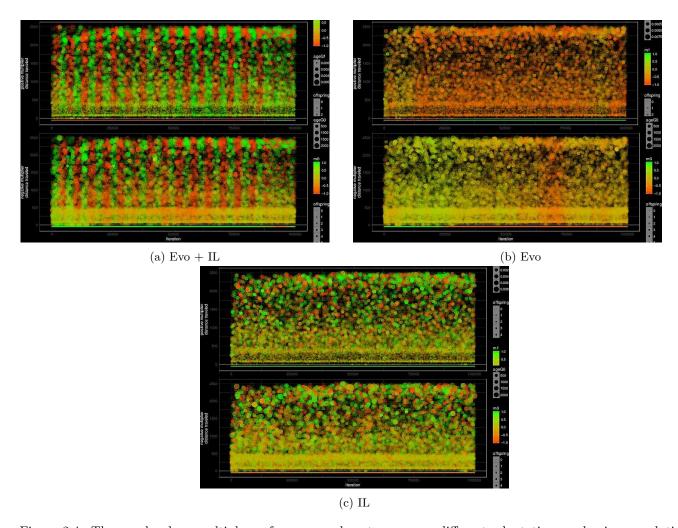


Figure 2.4: The graphs show multiple performance values to compare different adaptation mechanism: evolution combined with individual learning (\mathbf{a}), just evolution (\mathbf{b}) and just individual learning (\mathbf{c}). Data plotted shows 10% of the population (randomly sampled) and is taken from a single randomly selected run of the experiment. Each sub-graph shows the value of a single multiplier (names "positive" and "negative" refer to the initial value of the corresponding token), hence each genome has a data point on each of the sub-graphs. The values end-of-lifetime (x position), reproductive success (transparency; measured in number of offspring), adaptation success (size; measured in negative/positive token avoided/collected per life-time) and distance travelled (y-value) are the same in each sub-graph. The red/green lines below the y-axis is made up of dots, each representing the learning sign LS of the individuals: red=negative and green=positive. Note that LS is negated in the lower sub-graph.

	Evo	IL	Evo + IL	Baseline	Perfect
1	0.5093	0.5038	0.5031	0.5079	0.5056

Table 2.7: Median value of totalPucksRatio over the last 5000 iteration (N:30). Seasonal change every 30k iterations for configuration count=300 and value=1150.

	IL	Evo + IL	Baseline	Perfect
Evo	= 2.3e-01	$< 5.07 \mathrm{e} ext{-}09$	< 1.13 e-02	< 2.88e-03
IL		$< 2.47 \mathrm{e} ext{-}16$	$< 8.09 \mathrm{e} ext{-}07$	< 2.68e-06
Evo + IL			$> 2.19 \mathrm{e} ext{-}07$	> 1.66e-02
Baseline				= 1.18e-01

Table 2.8: Showing p-values of pairwise comparison of experiments for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Seasonal change every 30k iterations for configuration count=300 and value=1150. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

2.2 Seasons by Experiments. Count: 300 Value: 1150

2.2.1 Experiment: Evo, Count: 300 Value: 1150

	0	5k	15k	30k
1	0.5301	0.4995	0.4960	0.5038

Table 2.9: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment Evo for configuration count=300 and value=1150.

	5k	15k	30k
0	> 9.09e-69	$> 6.76\mathrm{e} ext{-}55$	> 1.14e-33
5k		$> 1.89 \mathrm{e} ext{-}02$	< 3.86e-06
15k			< 8.75 e-14

Table 2.10: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Evo for configuration count=300 and value=1150. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

2.2.2 Experiment: IL, Count: 300 Value: 1150

	0	5k	15k	30k
1	0.5034	0.5006	0.4973	0.5031

Table 2.11: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment IL for configuration count=300 and value=1150.

	5k	15k	30k
0	$> 2.54 \mathrm{e} ext{-}05$	> 6.93e-09	= 5.56e-01
5k		= 7.61e-02	$< 6.88 \mathrm{e} ext{-}05$
15k			< 8.5e-09

Table 2.12: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment IL for configuration count=300 and value=1150. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

2.2.3 Experiment: Evo + IL, Count: 300 Value: 1150

	0	5k	15k	30k
1	0.5306	0.5029	0.4981	0.5093

Table 2.13: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment Evo + IL for configuration count=300 and value=1150.

	5k	15k	30k
0	$> 2.82\mathrm{e} ext{-}35$	$> 1.37\mathrm{e} ext{-}31$	> 3.24e-16
5k		$> 1.7 \mathrm{e} ext{-}02$	$< 2.96 \mathbf{e}\text{-}11$
15k			< 1.14e-18

Table 2.14: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Evo + IL for configuration count=300 and value=1150. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

2.2.4 Experiment: Baseline, Count: 300 Value: 1150

	0	5k	15k	30k
1	0.5029	0.5013	0.4929	0.5056

Table 2.15: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment Baseline for configuration count=300 and value=1150.

Γ		5k	15k	30k
Г	0	= 6.66e-02	> 2.53e- 33	$< 6.54 \mathrm{e} ext{-}06$
Г	5k		> 9.38e-17	< 1.24 e-08
	15k			$< 1.78 \mathrm{e} ext{-}52$

Table 2.16: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Baseline for configuration count=300 and value=1150. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

2.2.5 Experiment: Perfect, Count: 300 Value: 1150

	0	5k	15k	30k
1	0.5624	0.5043	0.4877	0.5079

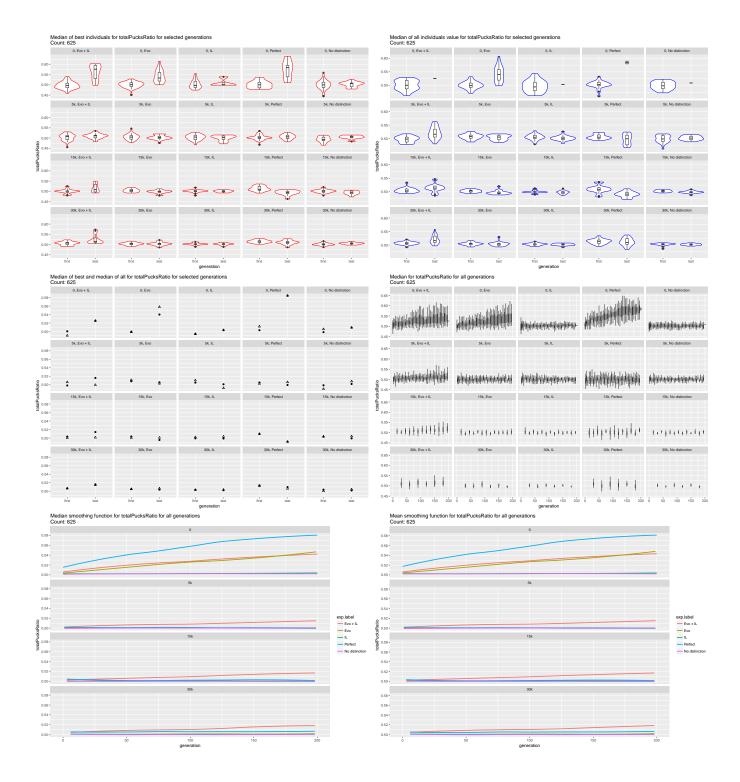
Table 2.17: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment Perfect for configuration count=300 and value=1150.

	5k	15k	30k
0	> 1.18e-101	> 8.48 e-85	> 2.59e-66
5k		> 1.49 e-12	< 5.78e- 05
15k			< 1.6e-33

Table 2.18: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Perfect for configuration count=300 and value=1150. The symbols $=, \neq, <, >$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

Chapter 3

Count: 625 Value: 625



3.1 Experiments by Seasons. Count: 625 Value: 625

3.1.1 Season: 0, Count: 625 Value: 625

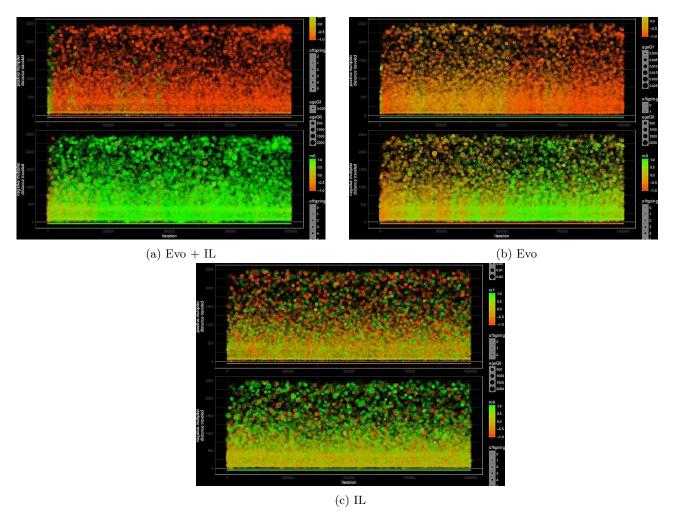


Figure 3.1: The graphs show multiple performance values to compare different adaptation mechanism: evolution combined with individual learning (\mathbf{a}), just evolution (\mathbf{b}) and just individual learning (\mathbf{c}). Data plotted shows 10% of the population (randomly sampled) and is taken from a single randomly selected run of the experiment. Each sub-graph shows the value of a single multiplier (names "positive" and "negative" refer to the initial value of the corresponding token), hence each genome has a data point on each of the sub-graphs. The values end-of-lifetime (x position), reproductive success (transparency; measured in number of offspring), adaptation success (size; measured in negative/positive token avoided/collected per life-time) and distance travelled (y-value) are the same in each sub-graph. The red/green lines below the y-axis is made up of dots, each representing the learning sign LS of the individuals: red=negative and green=positive. Note that LS is negated in the lower sub-graph.

		Evo	IL	Evo + IL	Baseline	Perfect
ĺ	1	0.5388	0.5411	0.5056	0.5868	0.5055

Table 3.1: Median value of totalPucksRatio over the last 5000 iteration (N:30). Seasonal change every 0 iterations for configuration count=625 and value=625.

	IL	Evo + IL	Baseline	Perfect
Evo	> 8.51e-77	= 4.32e-01	> 2.59e-74	$< 6.72 \mathrm{e} ext{-}52$
IL		< 6.66e- 59	= 1.31e-01	< 1.94 e-81
Evo + IL			> 5.52e- 59	$< 2.6 \mathrm{e} ext{-}34$
Baseline				< 1.88e-78

Table 3.2: Showing p-values of pairwise comparison of experiments for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Seasonal change every 0 iterations for configuration count=625 and value=625. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

3.1.2 Season: 5k, Count: 625 Value: 625

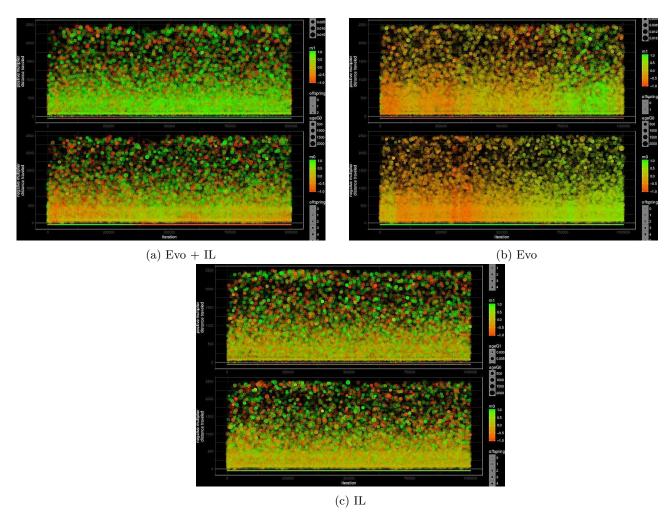


Figure 3.2: The graphs show multiple performance values to compare different adaptation mechanism: evolution combined with individual learning (\mathbf{a}), just evolution (\mathbf{b}) and just individual learning (\mathbf{c}). Data plotted shows 10% of the population (randomly sampled) and is taken from a single randomly selected run of the experiment. Each sub-graph shows the value of a single multiplier (names "positive" and "negative" refer to the initial value of the corresponding token), hence each genome has a data point on each of the sub-graphs. The values end-of-lifetime (x position), reproductive success (transparency; measured in number of offspring), adaptation success (size; measured in negative/positive token avoided/collected per life-time) and distance travelled (y-value) are the same in each sub-graph. The red/green lines below the y-axis is made up of dots, each representing the learning sign LS of the individuals: red=negative and green=positive. Note that LS is negated in the lower sub-graph.

	Evo	IL	Evo + IL	Baseline	Perfect
1	0.5134	0.4982	0.4975	0.4995	0.5004

Table 3.3: Median value of totalPucksRatio over the last 5000 iteration (N:30). Seasonal change every 5k iterations for configuration count=625 and value=625.

	IL	Evo + IL	Baseline	Perfect
Evo	> 4.42e-06	$< 3.36 \mathrm{e} ext{-}55$	= 1.18e-01	= 9.62e-01
IL		< 3.6e-70	< 1.72 e-10	< 2.41e-02
Evo + IL			> 3.6e-52	$> 1.09 \mathrm{e} ext{-}37$
Baseline				= 3.05e-01

Table 3.4: Showing p-values of pairwise comparison of experiments for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Seasonal change every 5k iterations for configuration count=625 and value=625. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

3.1.3 Season: 15k, Count: 625 Value: 625

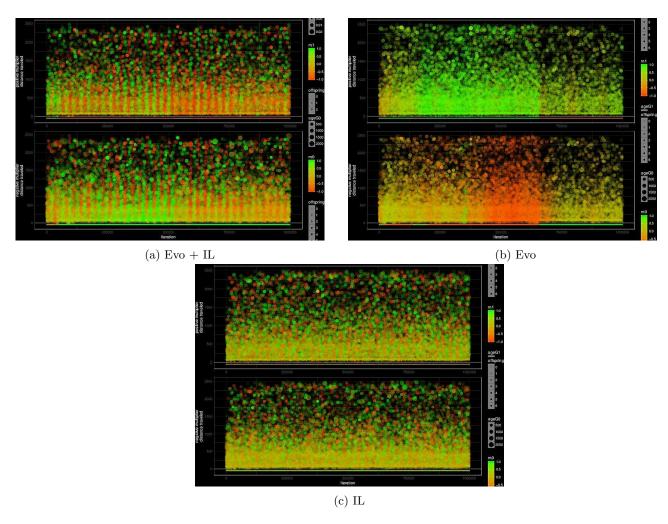


Figure 3.3: The graphs show multiple performance values to compare different adaptation mechanism: evolution combined with individual learning (\mathbf{a}), just evolution (\mathbf{b}) and just individual learning (\mathbf{c}). Data plotted shows 10% of the population (randomly sampled) and is taken from a single randomly selected run of the experiment. Each sub-graph shows the value of a single multiplier (names "positive" and "negative" refer to the initial value of the corresponding token), hence each genome has a data point on each of the sub-graphs. The values end-of-lifetime (x position), reproductive success (transparency; measured in number of offspring), adaptation success (size; measured in negative/positive token avoided/collected per life-time) and distance travelled (y-value) are the same in each sub-graph. The red/green lines below the y-axis is made up of dots, each representing the learning sign LS of the individuals: red=negative and green=positive. Note that LS is negated in the lower sub-graph.

	Evo	IL	Evo + IL	Baseline	Perfect
1	0.5136	0.4950	0.4993	0.4924	0.4990

Table 3.5: Median value of totalPucksRatio over the last 5000 iteration (N:30). Seasonal change every 15k iterations for configuration count=625 and value=625.

	IL	Evo + IL	Baseline	Perfect
Evo	< 7.4 e-03	< 1.09e-41	< 2.11e-07	> 1.07 e-07
IL		< 5.83e-42	= 2.01e-01	> 1.36 e-07
Evo + IL			> 1.04e-44	> 1.51e- 50
Baseline				> 3.37e-12

Table 3.6: Showing p-values of pairwise comparison of experiments for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Seasonal change every 15k iterations for configuration count=625 and value=625. The symbols $=, \neq, <, >$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

3.1.4 Season: 30k, Count: 625 Value: 625

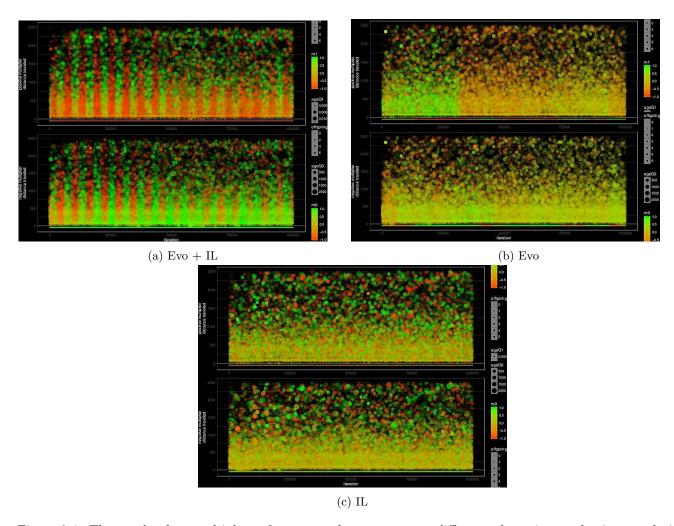


Figure 3.4: The graphs show multiple performance values to compare different adaptation mechanism: evolution combined with individual learning (\mathbf{a}), just evolution (\mathbf{b}) and just individual learning (\mathbf{c}). Data plotted shows 10% of the population (randomly sampled) and is taken from a single randomly selected run of the experiment. Each sub-graph shows the value of a single multiplier (names "positive" and "negative" refer to the initial value of the corresponding token), hence each genome has a data point on each of the sub-graphs. The values end-of-lifetime (x position), reproductive success (transparency; measured in number of offspring), adaptation success (size; measured in negative/positive token avoided/collected per life-time) and distance travelled (y-value) are the same in each sub-graph. The red/green lines below the y-axis is made up of dots, each representing the learning sign LS of the individuals: red=negative and green=positive. Note that LS is negated in the lower sub-graph.

	Evo	IL	Evo + IL	Baseline	Perfect
1	0.5159	0.5056	0.5012	0.5087	0.5015

Table 3.7: Median value of totalPucksRatio over the last 5000 iteration (N:30). Seasonal change every 30k iterations for configuration count=625 and value=625.

	IL	Evo + IL	Baseline	Perfect
Evo	> 2.33e-06	$< 8.79 \mathrm{e} ext{-}43$	> 2.82e- 08	< 1.01e-09
IL		< 1.33 e-65	= 9.15e-01	< 1.78e-21
Evo + IL			> 3.52e-74	> 8.02e-15
Baseline				< 2.77e-25

Table 3.8: Showing p-values of pairwise comparison of experiments for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Seasonal change every 30k iterations for configuration count=625 and value=625. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

3.2 Seasons by Experiments. Count: 625 Value: 625

3.2.1 Experiment: Evo, Count: 625 Value: 625

	0	5k	15k	30k
1	0.5411	0.4982	0.4950	0.5056

Table 3.9: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment Evo for configuration count=625 and value=625.

	5k	15k	30k
0	> 1.2e-131	> 4.34 e-99	> 7.43e-74
5k		$> 1.67 \mathrm{e}\text{-}05$	< 1.43e-12
15k			$< 1.35 \mathrm{e} ext{-}31$

Table 3.10: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Evo for configuration count=625 and value=625. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

3.2.2 Experiment: IL, Count: 625 Value: 625

		0	5k	15k	30k
Ì	1	0.5056	0.4975	0.4993	0.5012

Table 3.11: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment IL for configuration count=625 and value=625.

	5k	15k	30k
0	$>4.05\mathrm{e} ext{-}27$	>4.43e-20	$> 3.9 \mathrm{e} ext{-}06$
5k		= 8.51e-02	< 2.14e-16
15k			< 9.85e-14

Table 3.12: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment IL for configuration count=625 and value=625. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

3.2.3 Experiment: Evo + IL, Count: 625 Value: 625

	0	5k	15k	30k
1	0.5388	0.5134	0.5136	0.5159

Table 3.13: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment Evo + IL for configuration count=625 and value=625.

	5k	15k	30k
0	$> 1.76\mathrm{e} ext{-}32$	$>4.15\mathrm{e} ext{-}32$	> 7.06e-18
5k		=4.38e-01	$< 1.57 \mathrm{e} ext{-}07$
15k			< 6.91e-07

Table 3.14: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Evo + IL for configuration count=625 and value=625. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

3.2.4 Experiment: Baseline, Count: 625 Value: 625

	0	5k	15k	30k
1	0.5055	0.5004	0.4990	0.5015

Table 3.15: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment Baseline for configuration count=625 and value=625.

	5k	15k	30k
0	$> 3.21 \mathrm{e} ext{-}07$	>7.35e-18	$> 1.24 \mathrm{e} ext{-}05$
5k		> 4.61 e-04	$< 1.39 \mathrm{e} ext{-}02$
15k			$< 4.54 \mathrm{e} ext{-}15$

Table 3.16: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Baseline for configuration count=625 and value=625. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

3.2.5 Experiment: Perfect, Count: 625 Value: 625

	0	5k	15k	30k
1	0.5868	0.4995	0.4924	0.5087

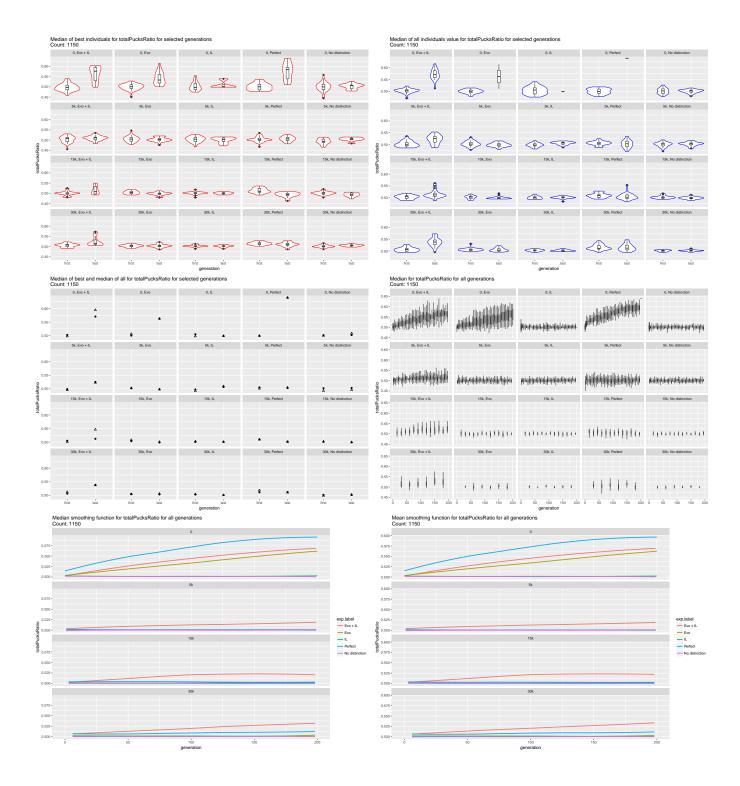
Table 3.17: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment Perfect for configuration count=625 and value=625.

	5k	15k	30k
0	> 1.58e-98	> 6.55e-82	> 2.09 e-76
5k		> 1.94 e-06	< 6.84 e-15
15k			< 2.06 e-35

Table 3.18: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Perfect for configuration count=625 and value=625. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

Chapter 4

Count: 1150 Value: 425



4.1 Experiments by Seasons. Count: 1150 Value: 425

4.1.1 Season: 0, Count: 1150 Value: 425

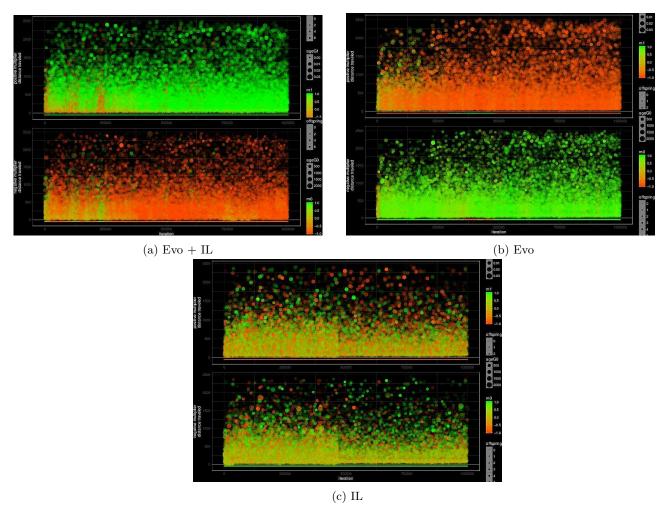


Figure 4.1: The graphs show multiple performance values to compare different adaptation mechanism: evolution combined with individual learning (\mathbf{a}), just evolution (\mathbf{b}) and just individual learning (\mathbf{c}). Data plotted shows 10% of the population (randomly sampled) and is taken from a single randomly selected run of the experiment. Each sub-graph shows the value of a single multiplier (names "positive" and "negative" refer to the initial value of the corresponding token), hence each genome has a data point on each of the sub-graphs. The values end-of-lifetime (x position), reproductive success (transparency; measured in number of offspring), adaptation success (size; measured in negative/positive token avoided/collected per life-time) and distance travelled (y-value) are the same in each sub-graph. The red/green lines below the y-axis is made up of dots, each representing the learning sign LS of the individuals: red=negative and green=positive. Note that LS is negated in the lower sub-graph.

		Evo	IL	Evo + IL	Baseline	Perfect
ĺ	1	0.5705	0.5662	0.4997	0.5982	0.5008

Table 4.1: Median value of totalPucksRatio over the last 5000 iteration (N:30). Seasonal change every 0 iterations for configuration count=1150 and value=425.

	IL	Evo + IL	Baseline	Perfect
Evo	> 4.44 e-82	< 1.5 e-03	> 1.51e-114	< 1.2e-51
IL		< 6.7e-150	= 3.43e-01	< 1.79e-111
Evo + IL			> 5.84 e-234	$< 2.95 \mathrm{e} ext{-}63$
Baseline				< 8.2e-152

Table 4.2: Showing p-values of pairwise comparison of experiments for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Seasonal change every 0 iterations for configuration count=1150 and value=425. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

4.1.2 Season: 5k, Count: 1150 Value: 425

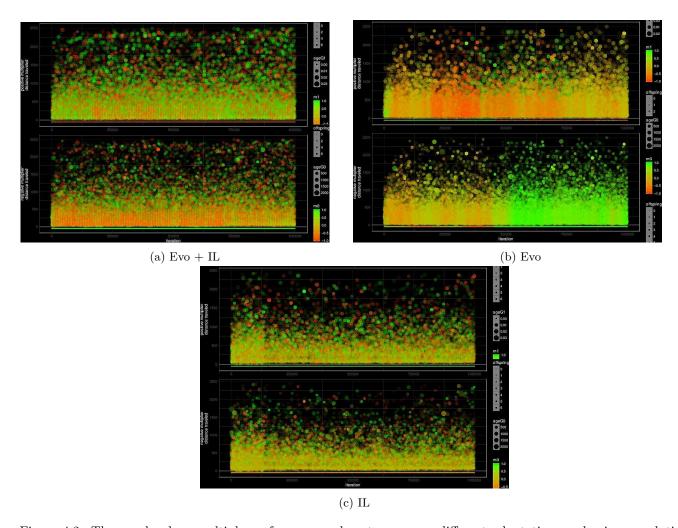


Figure 4.2: The graphs show multiple performance values to compare different adaptation mechanism: evolution combined with individual learning (\mathbf{a}), just evolution (\mathbf{b}) and just individual learning (\mathbf{c}). Data plotted shows 10% of the population (randomly sampled) and is taken from a single randomly selected run of the experiment. Each sub-graph shows the value of a single multiplier (names "positive" and "negative" refer to the initial value of the corresponding token), hence each genome has a data point on each of the sub-graphs. The values end-of-lifetime (x position), reproductive success (transparency; measured in number of offspring), adaptation success (size; measured in negative/positive token avoided/collected per life-time) and distance travelled (y-value) are the same in each sub-graph. The red/green lines below the y-axis is made up of dots, each representing the learning sign LS of the individuals: red=negative and green=positive. Note that LS is negated in the lower sub-graph.

	Evo	IL	Evo + IL	Baseline	Perfect
1	0.5191	0.4989	0.5023	0.5034	0.5003

Table 4.3: Median value of totalPucksRatio over the last 5000 iteration (N:30). Seasonal change every 5k iterations for configuration count=1150 and value=425.

	IL	Evo + IL	Baseline	Perfect
Evo	$< 4.22 \mathrm{e} ext{-}15$	$< 9.94 \mathrm{e} ext{-}122$	$< 5.77 \mathrm{e} ext{-}05$	< 3.73e-07
IL		< 2.9 e-80	> 2.92e-06	= 8.85e-01
Evo + IL			> 1.2e-102	$> 3.14 \mathrm{e} ext{-}72$
Baseline				< 5.28e-03

Table 4.4: Showing p-values of pairwise comparison of experiments for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Seasonal change every 5k iterations for configuration count=1150 and value=425. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

4.1.3 Season: 15k, Count: 1150 Value: 425

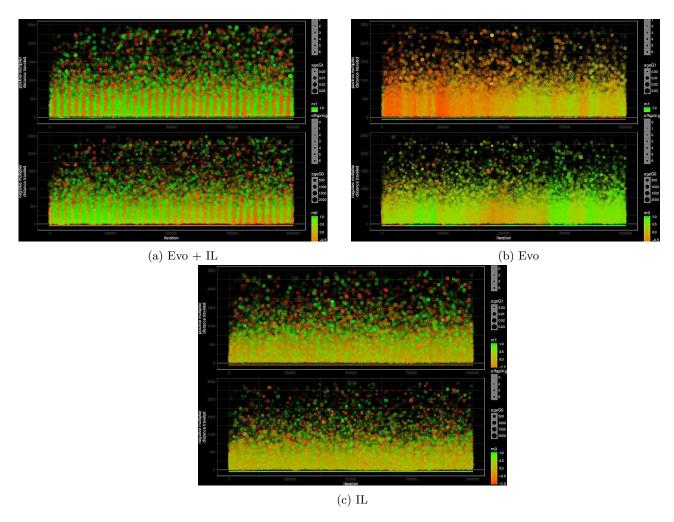


Figure 4.3: The graphs show multiple performance values to compare different adaptation mechanism: evolution combined with individual learning (\mathbf{a}), just evolution (\mathbf{b}) and just individual learning (\mathbf{c}). Data plotted shows 10% of the population (randomly sampled) and is taken from a single randomly selected run of the experiment. Each sub-graph shows the value of a single multiplier (names "positive" and "negative" refer to the initial value of the corresponding token), hence each genome has a data point on each of the sub-graphs. The values end-of-lifetime (\mathbf{x} position), reproductive success (transparency; measured in number of offspring), adaptation success (size; measured in negative/positive token avoided/collected per life-time) and distance travelled (\mathbf{y} -value) are the same in each sub-graph. The red/green lines below the \mathbf{y} -axis is made up of dots, each representing the learning sign LS of the individuals: red=negative and green=positive. Note that LS is negated in the lower sub-graph.

	Evo	IL	Evo + IL	Baseline	Perfect
1	0.5121	0.4981	0.5011	0.5032	0.5001

Table 4.5: Median value of totalPucksRatio over the last 5000 iteration (N:30). Seasonal change every 15k iterations for configuration count=1150 and value=425.

	IL	Evo + IL	Baseline	Perfect
Evo	< 1.28 e-08	< 3.11e-77	= 5.2e-01	< 4.33e- 16
IL		< 2.23 e-72	> 2.2e-10	$< 1.97 \mathrm{e} ext{-}06$
Evo + IL			> 1.84e- 91	> 2.24e-28
Baseline				< 6.09e-20

Table 4.6: Showing p-values of pairwise comparison of experiments for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Seasonal change every 15k iterations for configuration count=1150 and value=425. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

4.1.4 Season: 30k, Count: 1150 Value: 425

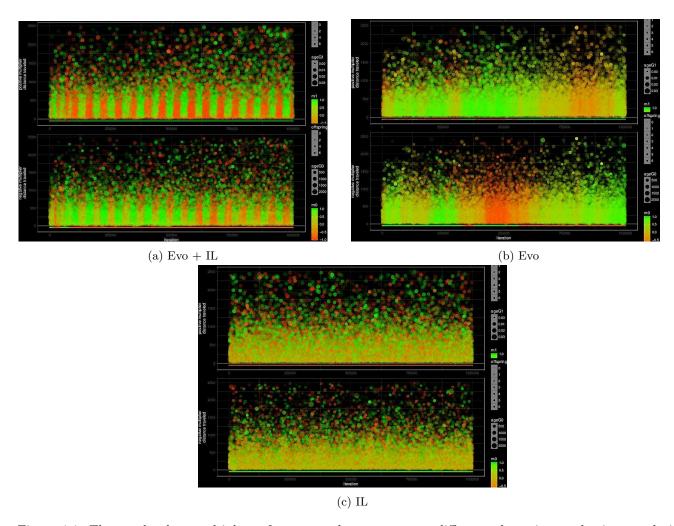


Figure 4.4: The graphs show multiple performance values to compare different adaptation mechanism: evolution combined with individual learning (\mathbf{a}), just evolution (\mathbf{b}) and just individual learning (\mathbf{c}). Data plotted shows 10% of the population (randomly sampled) and is taken from a single randomly selected run of the experiment. Each sub-graph shows the value of a single multiplier (names "positive" and "negative" refer to the initial value of the corresponding token), hence each genome has a data point on each of the sub-graphs. The values end-of-lifetime (x position), reproductive success (transparency; measured in number of offspring), adaptation success (size; measured in negative/positive token avoided/collected per life-time) and distance travelled (y-value) are the same in each sub-graph. The red/green lines below the y-axis is made up of dots, each representing the learning sign LS of the individuals: red=negative and green=positive. Note that LS is negated in the lower sub-graph.

	Evo	IL	Evo + IL	Baseline	Perfect
1	0.5334	0.5032	0.5003	0.5130	0.5003

Table 4.7: Median value of totalPucksRatio over the last 5000 iteration (N:30). Seasonal change every 30k iterations for configuration count=1150 and value=425.

	IL	Evo + IL	Baseline	Perfect
Evo	> 8.82 e-14	< 7.94e-74	> 9.45e-15	< 3.28e-33
IL		< 7.58e-82	= 9.59e-01	< 4.58e-64
Evo + IL			> 5.74 e-77	> 1.08e-27
Baseline				< 1.41e-58

Table 4.8: Showing p-values of pairwise comparison of experiments for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Seasonal change every 30k iterations for configuration count=1150 and value=425. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

4.2 Seasons by Experiments. Count: 1150 Value: 425

4.2.1 Experiment: Evo, Count: 1150 Value: 425

	0	5k	15k	30k
1	0.5662	0.4989	0.4981	0.5032

Table 4.9: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment Evo for configuration count=1150 and value=425.

	5k	15k	30k
0	$> 6.91 \mathrm{e} ext{-}120$	> 9.94e-88	> 5.28e- 79
5k		< 6.04 e-03	< 5.82 e-36
15k			< 1.14e-27

Table 4.10: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Evo for configuration count=1150 and value=425. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

4.2.2 Experiment: IL, Count: 1150 Value: 425

	0	5k	15k	30k
1	0.4997	0.5023	0.5011	0.5003

Table 4.11: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment IL for configuration count=1150 and value=425.

	5k	15k	30k
0	= 1.03e-01	= 7.08e-01	= 3.65e-01
5k		> 8.33e- 03	> 3e-02
15k			= 5.26e-01

Table 4.12: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment IL for configuration count=1150 and value=425. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

4.2.3 Experiment: Evo + IL, Count: 1150 Value: 425

	0	5k	15k	30k
1	0.5705	0.5191	0.5121	0.5334

Table 4.13: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment Evo + IL for configuration count=1150 and value=425.

	5k	15k	30k
0	$> 3.41 \mathrm{e} ext{-}178$	$> 3.54 \mathrm{e} ext{-}118$	$> 2.16\mathrm{e} ext{-}85$
5k		= 7.19e-01	< 1.87 e-25
15k			< 3.75 e-22

Table 4.14: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Evo + IL for configuration count=1150 and value=425. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

4.2.4 Experiment: Baseline, Count: 1150 Value: 425

	0	5k	15k	30k
1	0.5008	0.5003	0.5001	0.5003

Table 4.15: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment Baseline for configuration count=1150 and value=425.

	5k	15k	30k
0	= 2.33e-01	> 1.53e-06	= 8.36e-01
5k		> 5.33e- 03	= 1.6e-01
15k			< 2.24 e-08

Table 4.16: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Baseline for configuration count=1150 and value=425. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

4.2.5 Experiment: Perfect, Count: 1150 Value: 425

	0	5k	15k	30k
1	0.5982	0.5034	0.5032	0.5130

Table 4.17: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment Perfect for configuration count=1150 and value=425.

	5k	15k	30k
0	> 6e-153	> 3.08e-119	> 2.33e-124
5k		< 1.14e-04	< 2.97 e-53
15k			< 1.93e-28

Table 4.18: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Perfect for configuration count=1150 and value=425. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

Chapter 5

Count by Experiments. Season: 0 -

5.1 Season: 0 -

5.1.1 Experiment: Evo, Season: 0 -

	300	625	1150
1	0.5301	0.5411	0.5662

Table 5.1: Median value of *totalPucksRatio* over the last 5000 iteration (N:30). Experiment *Evo* for seasonal change every 0 iterations.

	count:625 value:625	count:1150 value:425
count:300 value:1150	$< 1.24 \mathrm{e} ext{-}07$	< 3.02e-27
count:625 value:625		< 9.37e-16

Table 5.2: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Evo for seasonal change every 0 iterations. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

5.1.2 Experiment: IL, Season: 0 -

	300	625	1150
1	0.5034	0.5056	0.4997

Table 5.3: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment IL for seasonal change every 0 iterations.

	count:625 value:625	count:1150 value:425
count:300 value:1150	= 5.78e-01	> 7.33e- 05
count:625 value:625		> 7.87e-11

Table 5.4: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment IL for seasonal change every 0 iterations. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

5.1.3 Experiment: Evo + IL, Season: 0 -

	300	625	1150
1	0.5306	0.5388	0.5705

Table 5.5: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment Evo + IL for seasonal change every 0 iterations.

	count:625 value:625	count:1150 value:425
count:300 value:1150	$< 3.27 \mathrm{e} ext{-}02$	< 1.44e-40
count:625 value:625		< 1.33e-32

Table 5.6: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Evo + IL for seasonal change every 0 iterations. The symbols $=, \neq, <, >$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

5.1.4 Experiment: Baseline, Season: 0 -

	300	625	1150
1	0.5029	0.5055	0.5008

Table 5.7: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment Baseline for seasonal change every 0 iterations.

	count:625 value:625	count:1150 value:425
count:300 value:1150	= 2.53e-01	> 8.47 e-07
count:625 value:625		> 3.72 e-11

Table 5.8: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment *Baseline* for seasonal change every 0 iterations. The symbols $=, \neq, <, >$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

5.1.5 Experiment: Perfect, Season: 0 -

	300	625	1150
1	0.5624	0.5868	0.5982

Table 5.9: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment Perfect for seasonal change every 0 iterations.

	count:625 value:625	count:1150 value:425
count:300 value:1150	< 3.81e-13	< 2.8e-44
count:625 value:625		< 1.91e-09

Table 5.10: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Perfect for seasonal change every 0 iterations. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

5.2 Season: 5k -

5.2.1 Experiment: Evo, Season: 5k -

	300	625	1150
1	0.4995	0.4982	0.4989

Table 5.11: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment Evo for seasonal change every 5k iterations.

	count:625 value:625	count:1150 value:425
count:300 value:1150	= 4.55e-01	= 3.08e-01
count:625 value:625		> 1.22e- 03

Table 5.12: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Evo for seasonal change every 5k iterations. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

5.2.2 Experiment: IL, Season: 5k -

	300	625	1150
1	0.5006	0.4975	0.5023

Table 5.13: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment IL for seasonal change every 5k iterations.

	count:625 value:625	count:1150 value:425
count:300 value:1150	> 7.89 e-05	< 1.99 e-02
count:625 value:625		< 1.81e-17

Table 5.14: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment IL for seasonal change every 5k iterations. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

5.2.3 Experiment: Evo + IL, Season: 5k -

	300	625	1150
1	0.5029	0.5134	0.5191

Table 5.15: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment Evo + IL for seasonal change every 5k iterations.

	count:625 value:625	count:1150 value:425
count:300 value:1150	$< 3.87 \mathrm{e} ext{-}19$	< 1.5 e-32
count:625 value:625		< 5.88e-04

Table 5.16: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Evo + IL for seasonal change every 5k iterations. The symbols $=, \neq, <, >$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

5.2.4 Experiment: Baseline, Season: 5k -

	300	625	1150
1	0.5013	0.5004	0.5003

Table 5.17: Median value of *totalPucksRatio* over the last 5000 iteration (N:30). Experiment *Baseline* for seasonal change every 5k iterations.

	count:625 value:625	count:1150 value:425
count:300 value:1150	= 5.98e-01	$>4.72\mathrm{e} ext{-}02$
count:625 value:625		$> 1.27 \mathrm{e} ext{-}02$

Table 5.18: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Baseline for seasonal change every 5k iterations. The symbols $=, \neq, <, >$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

5.2.5 Experiment: Perfect, Season: 5k -

	300	625	1150
1	0.5043	0.4995	0.5034

Table 5.19: Median value of *totalPucksRatio* over the last 5000 iteration (N:30). Experiment *Perfect* for seasonal change every 5k iterations.

	count:625 value:625	count:1150 value:425
count:300 value:1150	$>4.23\mathrm{e} ext{-}02$	= 2.01e-01
count:625 value:625		= 7.76e-02

Table 5.20: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Perfect for seasonal change every 5k iterations. The symbols $=, \neq, <, >$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

5.3 Season: 15k -

5.3.1 Experiment: Evo, Season: 15k -

	300	625	1150
1	0.4960	0.4950	0.4981

Table 5.21: Median value of *totalPucksRatio* over the last 5000 iteration (N:30). Experiment *Evo* for seasonal change every 15k iterations.

	count:625 value:625	count:1150 value:425
count:300 value:1150	$< 4.78 \mathrm{e} ext{-}02$	$< 1.58 \mathrm{e} ext{-}04$
count:625 value:625		< 1.09e-08

Table 5.22: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Evo for seasonal change every 15k iterations. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

5.3.2 Experiment: IL, Season: 15k -

	300	625	1150
1	0.4973	0.4993	0.5011

Table 5.23: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment IL for seasonal change every 15k iterations.

	count:625 value:625	count:1150 value:425
count:300 value:1150	= 6.51e-01	< 4.11e-04
count:625 value:625		< 1.44e-12

Table 5.24: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment IL for seasonal change every 15k iterations. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

5.3.3 Experiment: Evo + IL, Season: 15k -

	300	625	1150
1	0.4981	0.5136	0.5121

Table 5.25: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment Evo + IL for seasonal change every 15k iterations.

	count:625 value:625	count:1150 value:425
count:300 value:1150	< 1.94e- 16	< 9.56e- 30
count:625 value:625		= 2.61e-01

Table 5.26: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Evo + IL for seasonal change every 15k iterations. The symbols $=, \neq, <, >$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

5.3.4 Experiment: Baseline, Season: 15k -

	300	625	1150
1	0.4929	0.4990	0.5001

Table 5.27: Median value of *totalPucksRatio* over the last 5000 iteration (N:30). Experiment *Baseline* for seasonal change every 15k iterations.

	count:625 value:625	count:1150 value:425
count:300 value:1150	$< 6.05 \mathrm{e} ext{-}28$	$< 6.35 \mathrm{e} ext{-}27$
count:625 value:625		= 6.21e-01

Table 5.28: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Baseline for seasonal change every 15k iterations. The symbols $=, \neq, <, >$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

5.3.5 Experiment: Perfect, Season: 15k -

	300	625	1150
1	0.4877	0.4924	0.5032

Table 5.29: Median value of *totalPucksRatio* over the last 5000 iteration (N:30). Experiment *Perfect* for seasonal change every 15k iterations.

	count:625 value:625	count:1150 value:425
count:300 value:1150	< 1.08 e-03	$< 6.24\mathrm{e} ext{-}31$
count:625 value:625		< 2.41e-28

Table 5.30: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Perfect for seasonal change every 15k iterations. The symbols $=, \neq, <, >$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

5.4 Season: 30k -

5.4.1 Experiment: Evo, Season: 30k -

	300	625	1150
1	0.5038	0.5056	0.5032

Table 5.31: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment Evo for seasonal change every 30k iterations.

	count:625 value:625	count:1150 value:425
count:300 value:1150	= 6.73e-01	= 4.04e-01
count:625 value:625		= 8.4e-01

Table 5.32: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Evo for seasonal change every 30k iterations. The symbols $=, \neq, <, >$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

5.4.2 Experiment: IL, Season: 30k -

	300	625	1150
1	0.5031	0.5012	0.5003

Table 5.33: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment IL for seasonal change every 30k iterations.

	count:625 value:625	count:1150 value:425
count:300 value:1150	> 1.87 e-04	> 4.62e-09
count:625 value:625		> 3.55e- 02

Table 5.34: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment IL for seasonal change every 30k iterations. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

5.4.3 Experiment: Evo + IL, Season: 30k -

	300	625	1150
1	0.5093	0.5159	0.5334

Table 5.35: Median value of totalPucksRatio over the last 5000 iteration (N:30). Experiment Evo + IL for seasonal change every 30k iterations.

	count:625 value:625	count:1150 value:425
count:300 value:1150	< 4.7e-09	$< \mathbf{2.64e\text{-}31}$
count:625 value:625		$< 3.59 \mathrm{e} ext{-}14$

Table 5.36: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Evo + IL for seasonal change every 30k iterations. The symbols $=, \neq, <, >$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

5.4.4 Experiment: Baseline, Season: 30k -

	300	625	1150
1	0.5056	0.5015	0.5003

Table 5.37: Median value of *totalPucksRatio* over the last 5000 iteration (N:30). Experiment *Baseline* for seasonal change every 30k iterations.

	count:625 value:625	count:1150 value:425
count:300 value:1150	$>4.77\mathrm{e} ext{-}33$	$>9.37\mathrm{e} ext{-}42$
count:625 value:625		> 4.81e-03

Table 5.38: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Baseline for seasonal change every 30k iterations. The symbols $=, \neq, <, >$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.

5.4.5 Experiment: Perfect, Season: 30k -

	300	625	1150
1	0.5079	0.5087	0.5130

Table 5.39: Median value of *totalPucksRatio* over the last 5000 iteration (N:30). Experiment *Perfect* for seasonal change every 30k iterations.

	count:625 value:625	count:1150 value:425
count:300 value:1150	= 1.6e-01	$< 6.76 \mathrm{e} ext{-}09$
count:625 value:625		< 3.43e-06

Table 5.40: Showing p-values of pairwise comparison of seasonal change for totalPucksRatio (row vs. column) over the last 5000 iteration (N:30). Experiment Perfect for seasonal change every 30k iterations. The symbols $=,\neq,<,>$, indicate whether the median values for totalPucksRatio are equal, not equal (but not significantly larger or smaller), significantly smaller or significantly larger respectively. p-values below the significance level of 0.05 are written in bold.