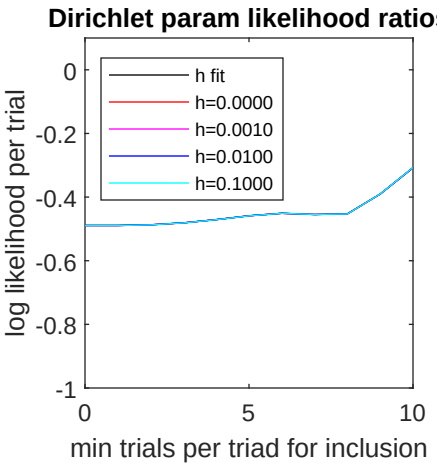

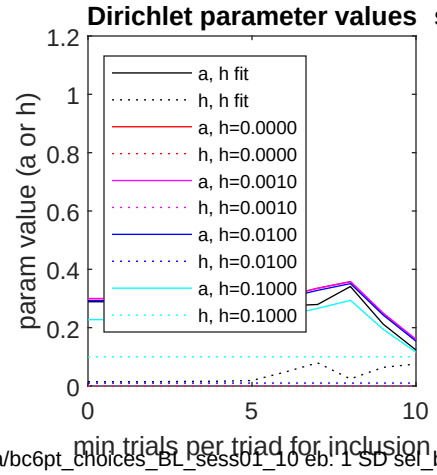

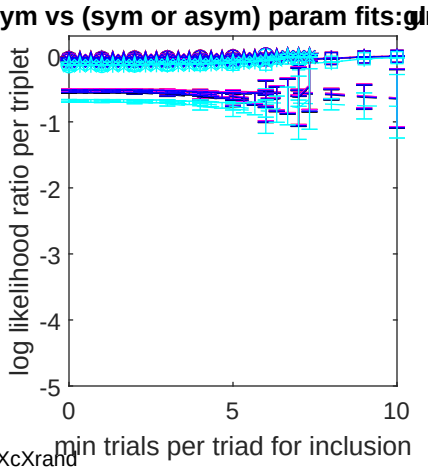

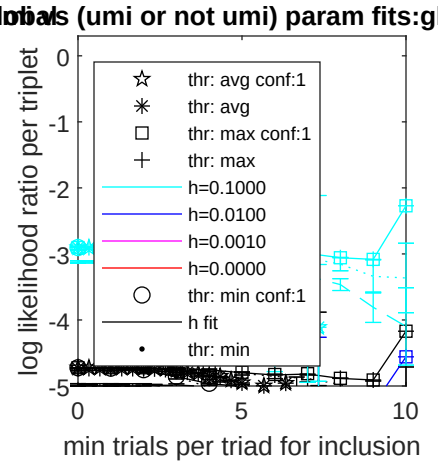

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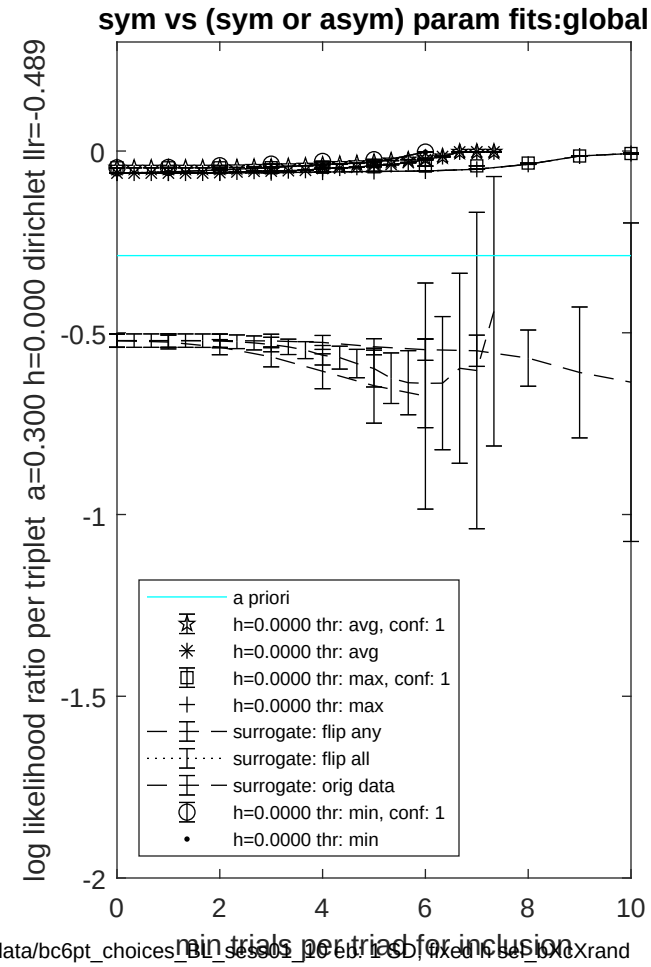

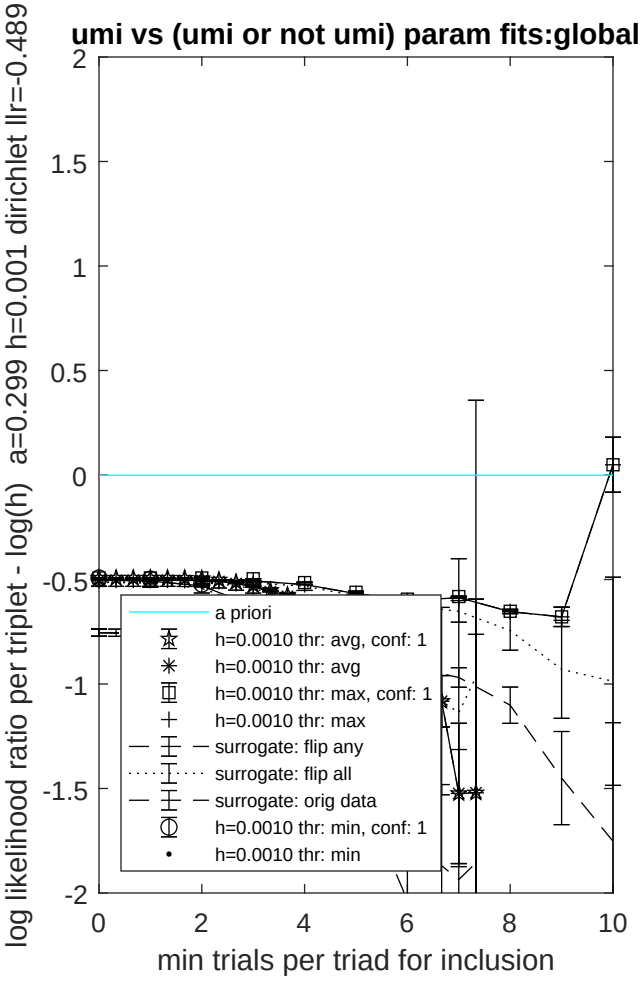

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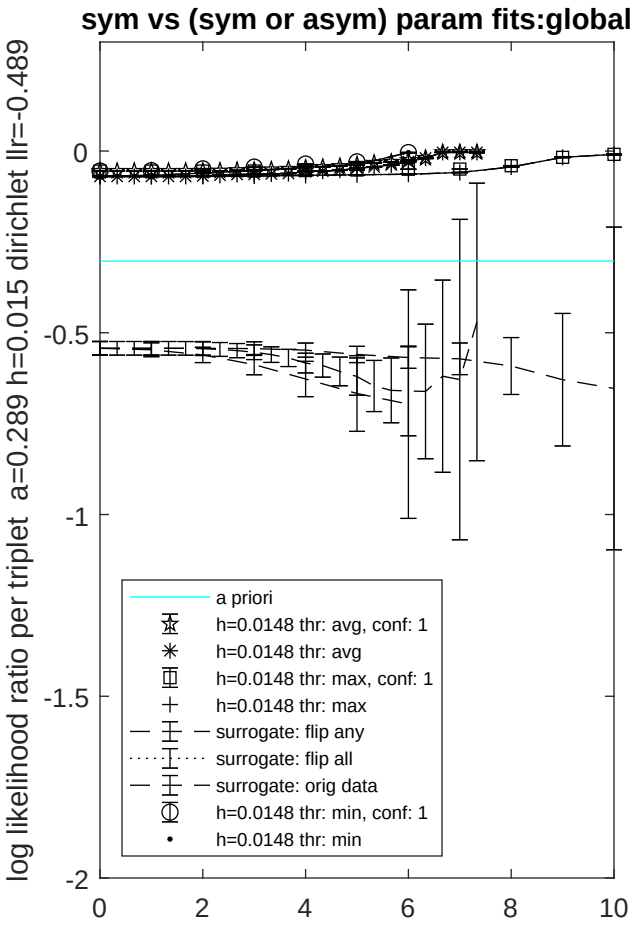

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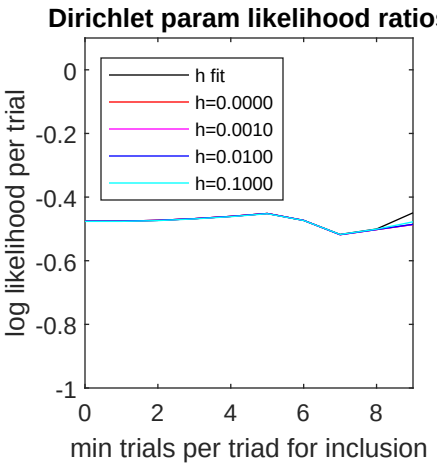

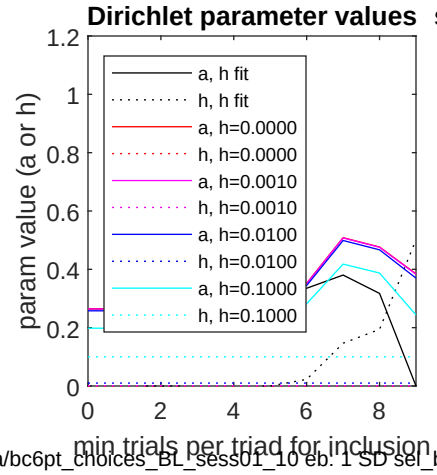

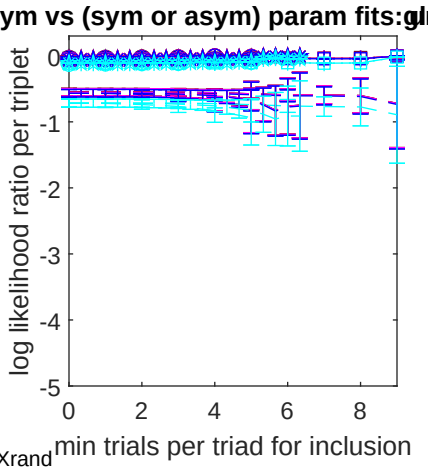

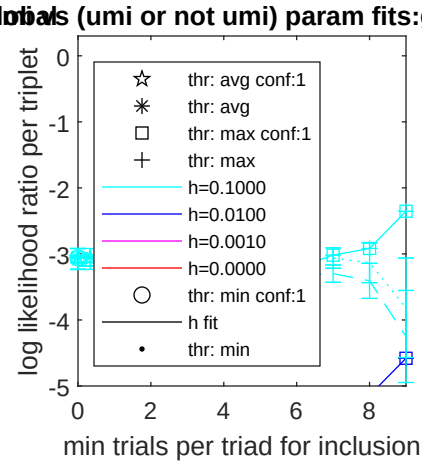

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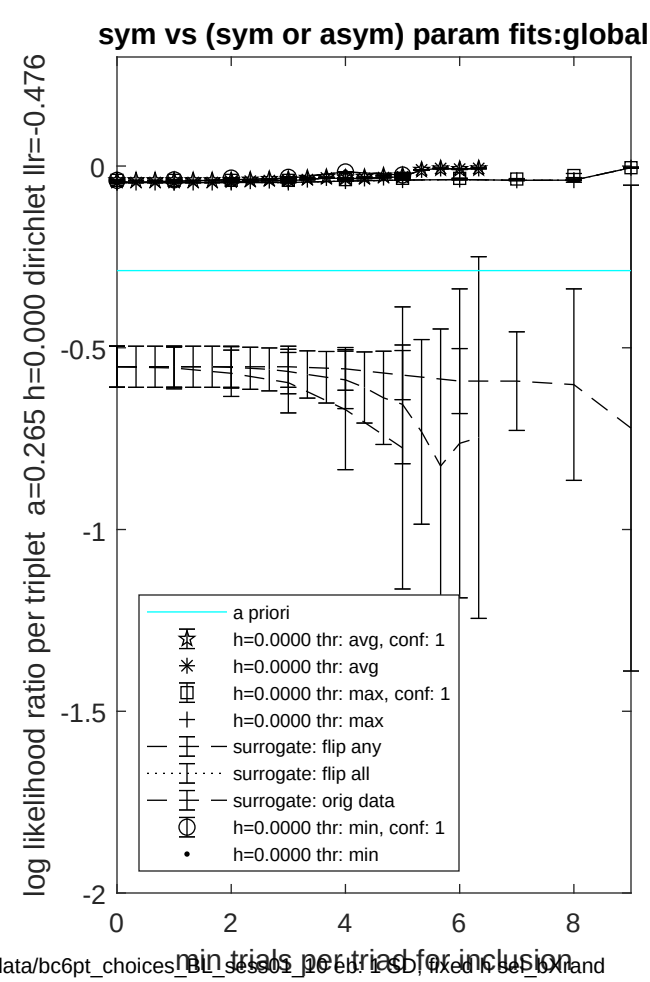

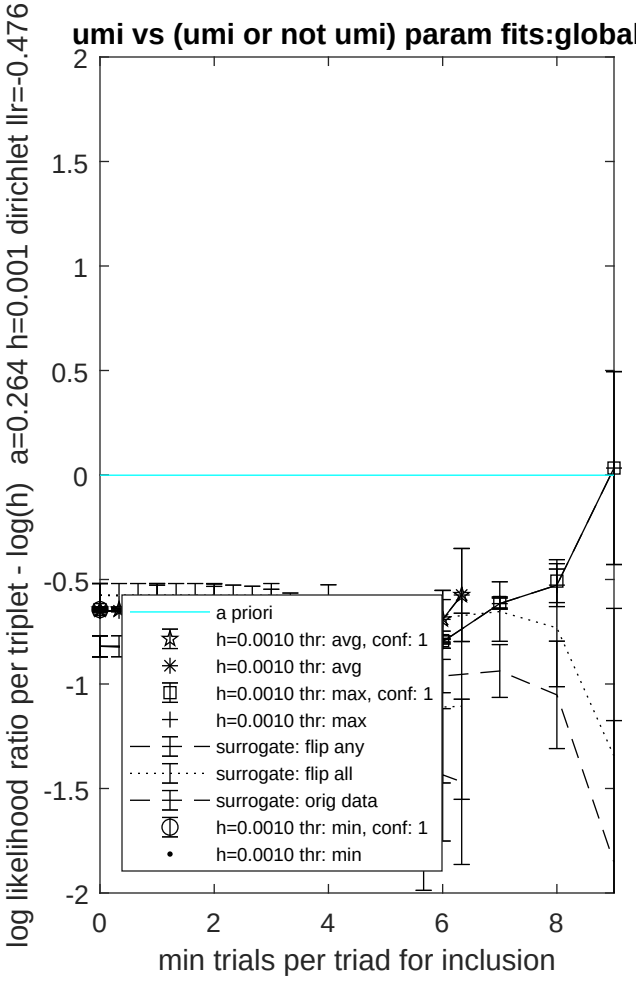

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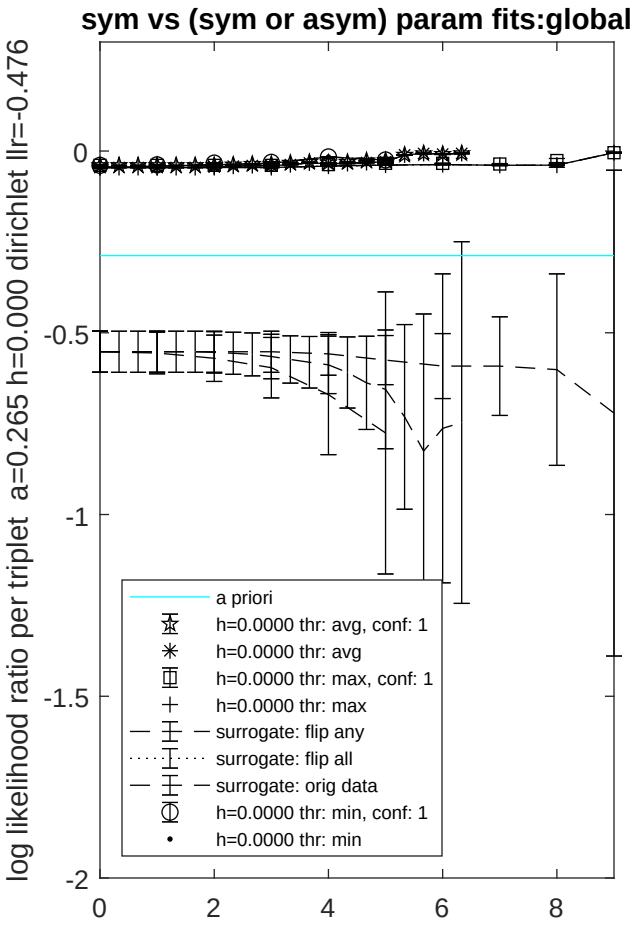

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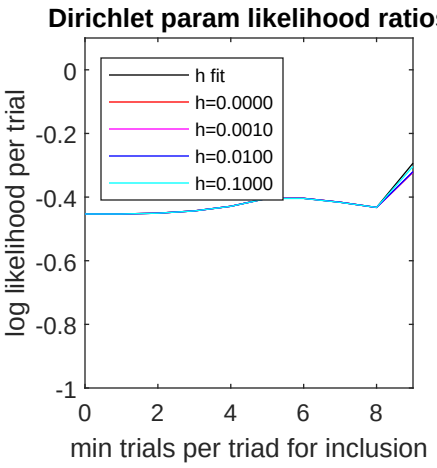

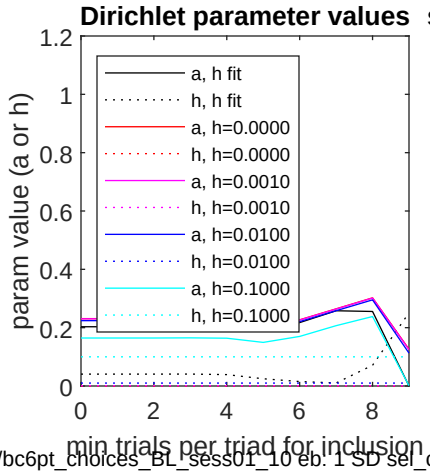

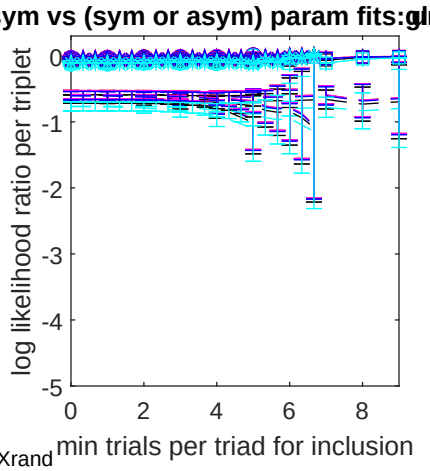

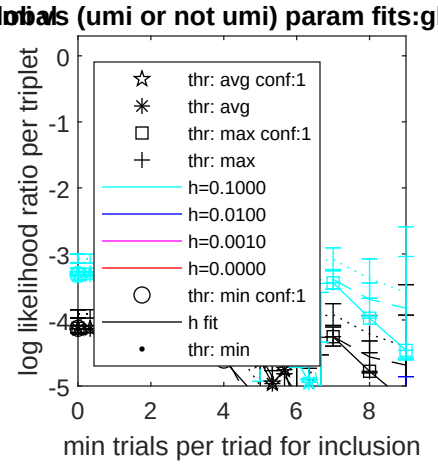

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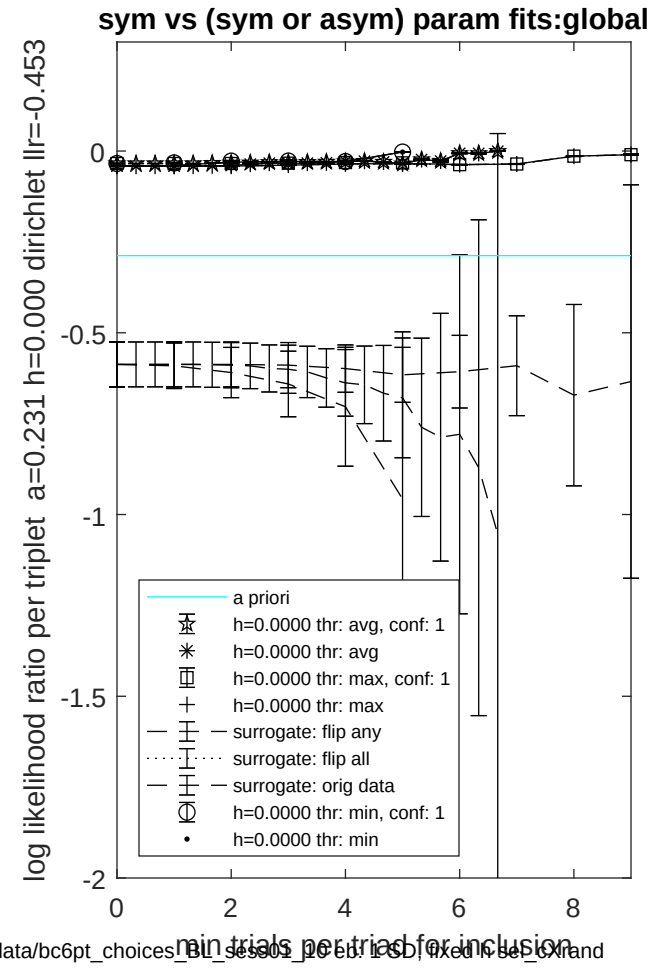

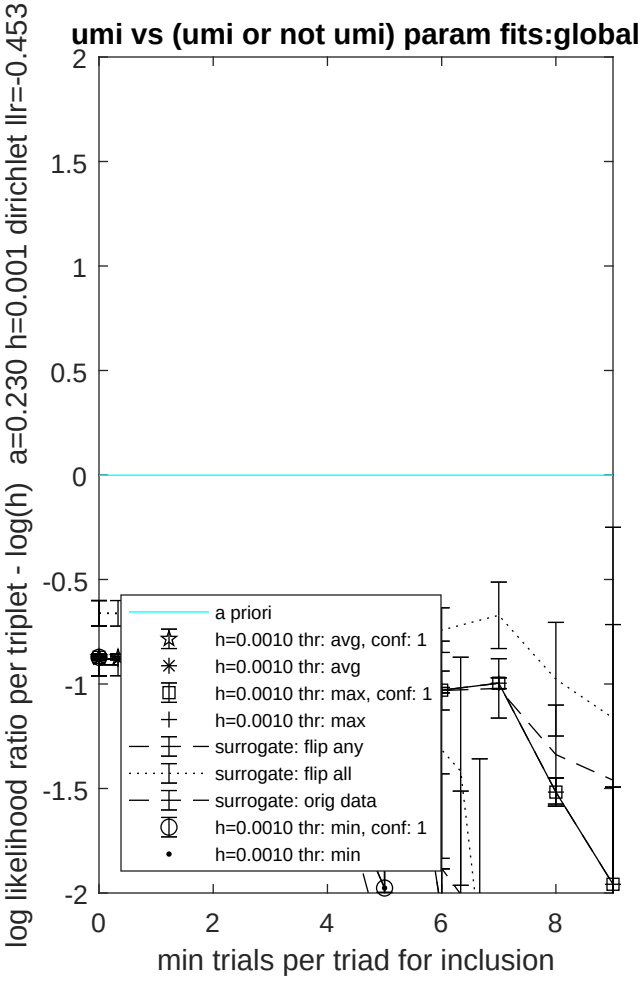

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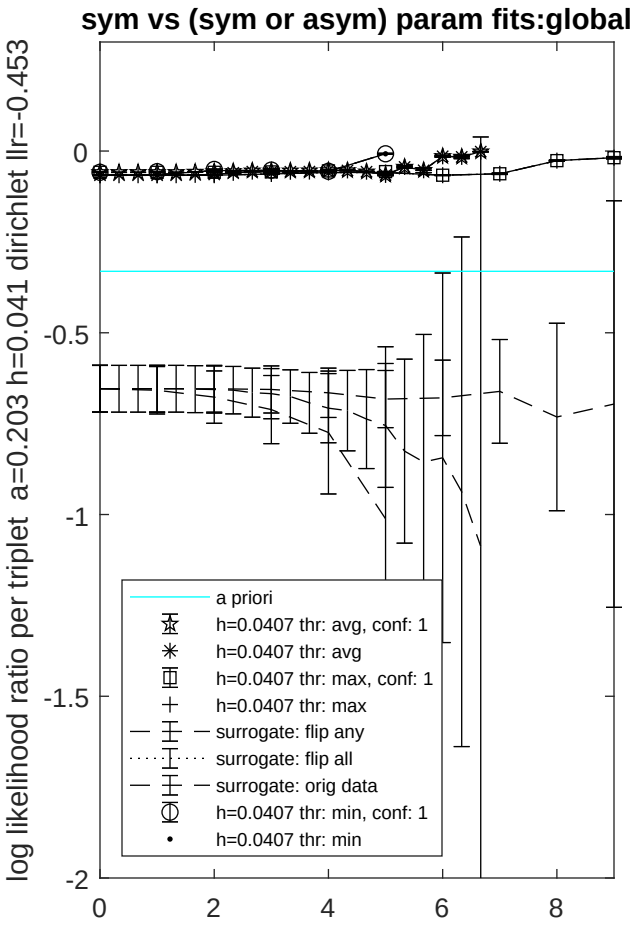

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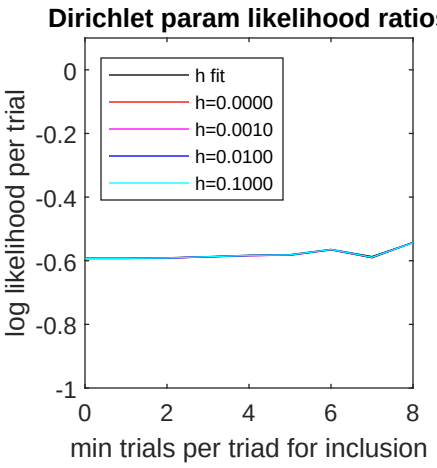

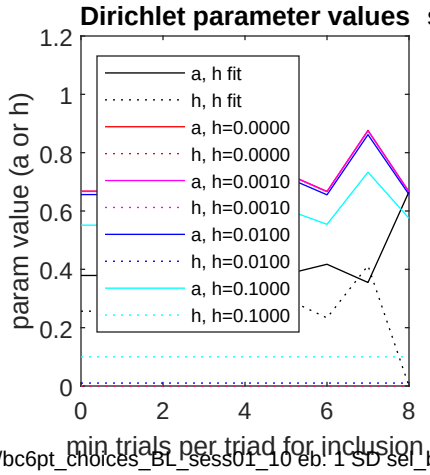

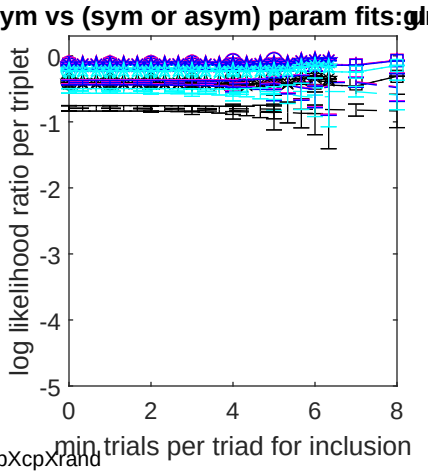

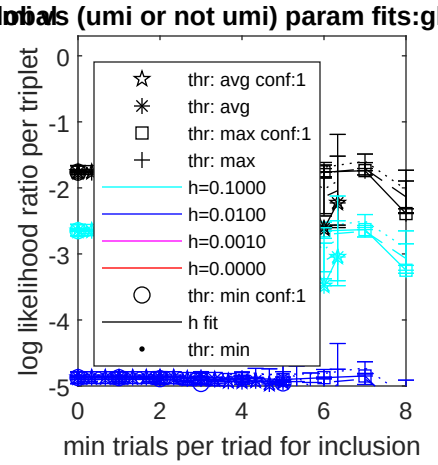

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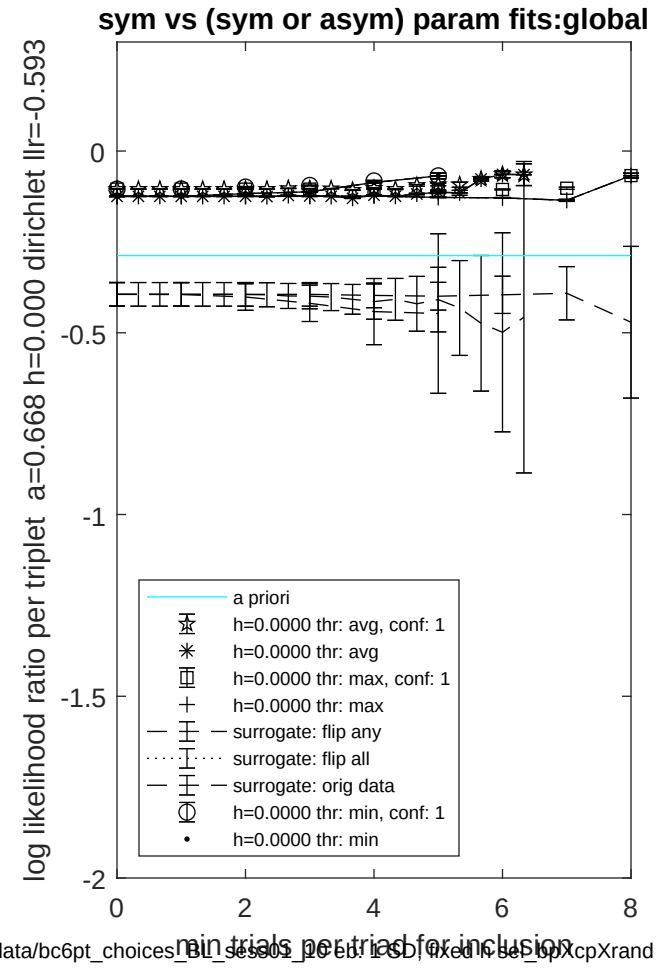

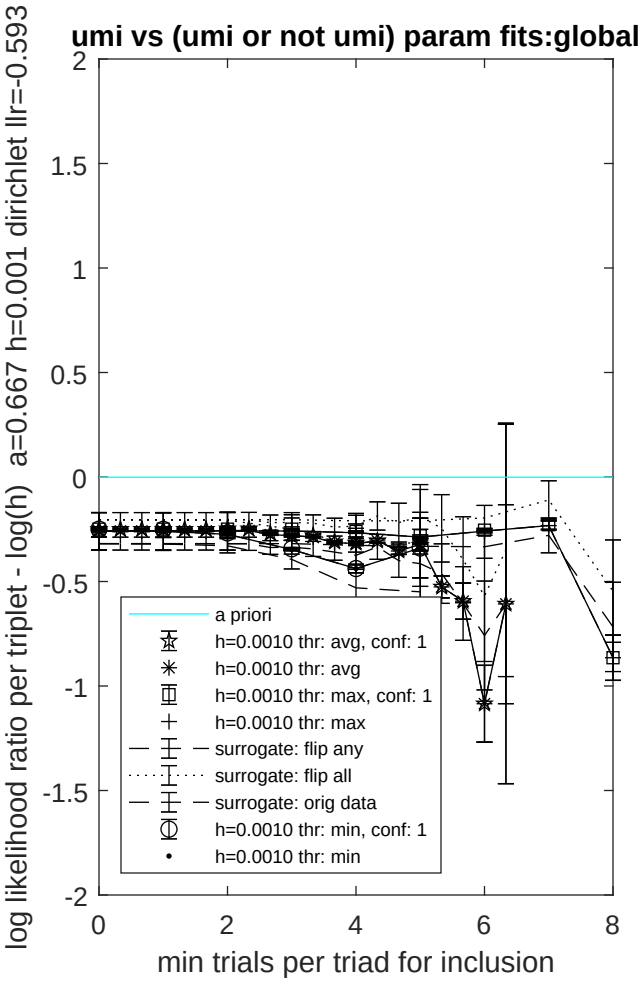

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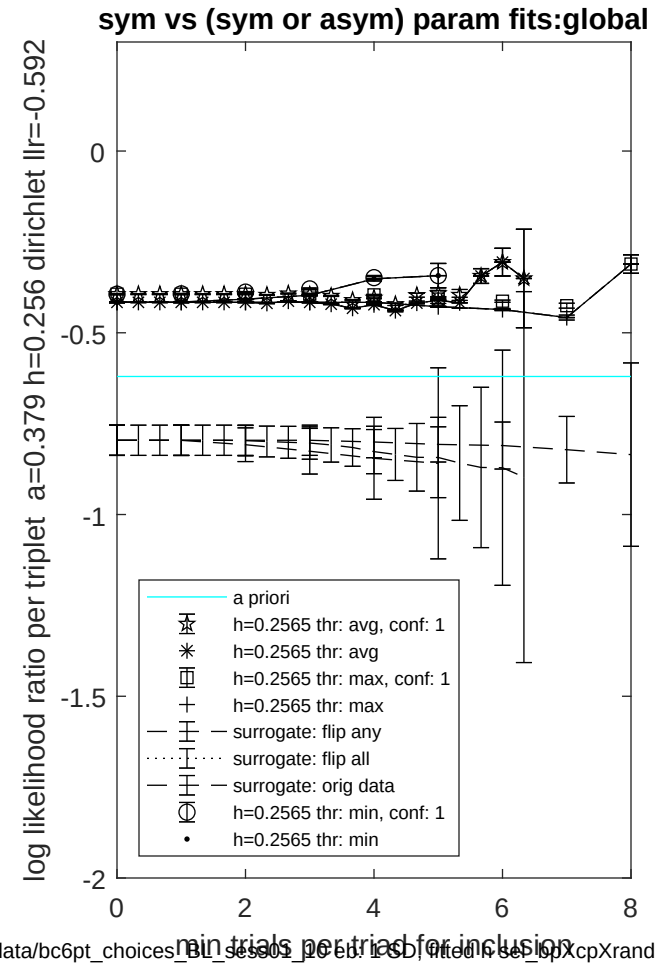

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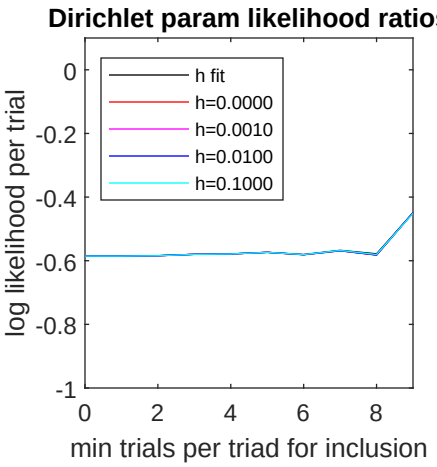

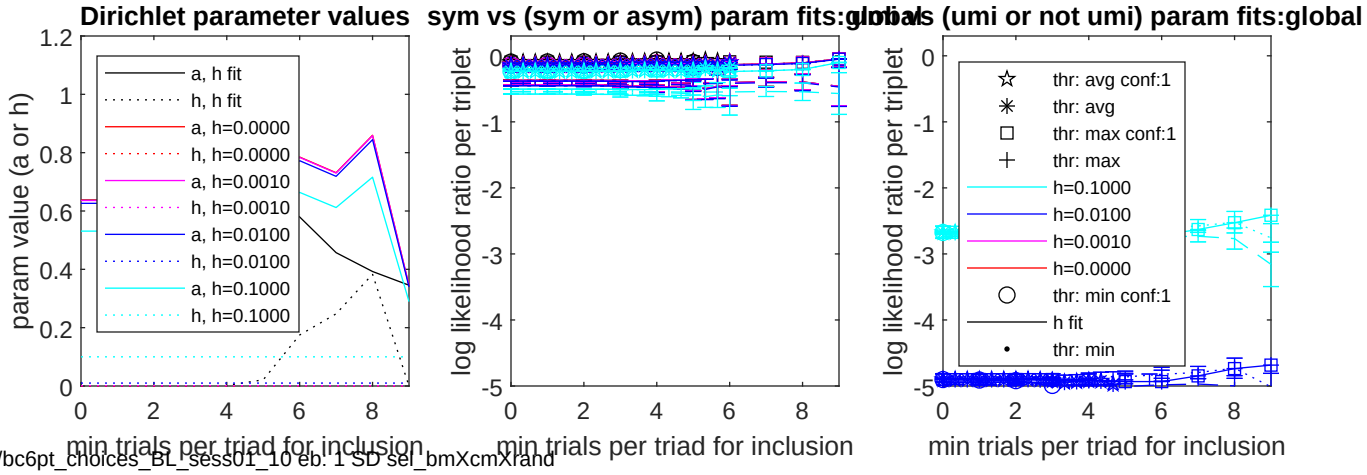

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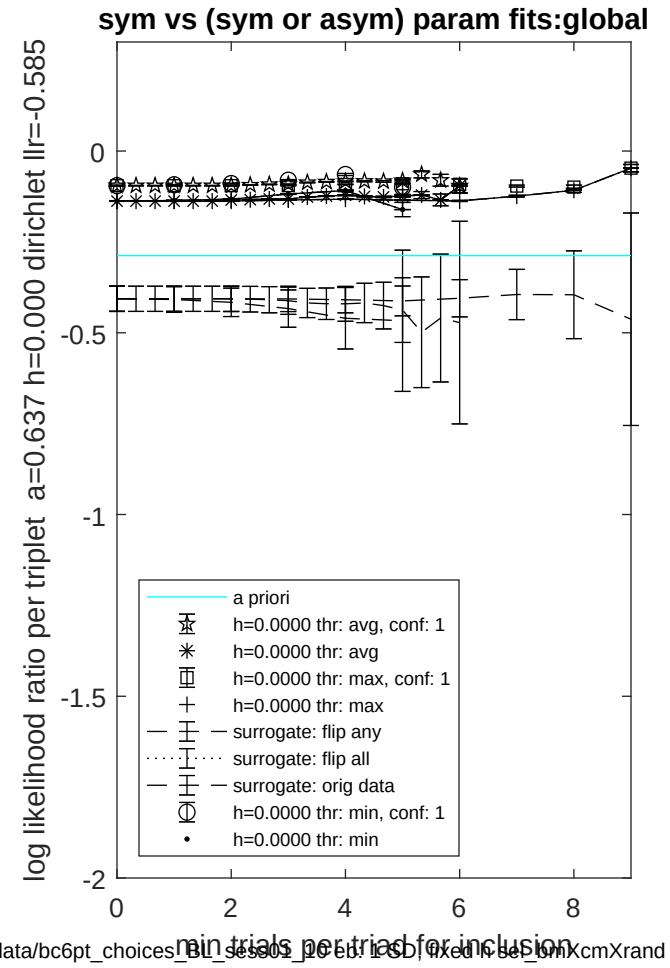

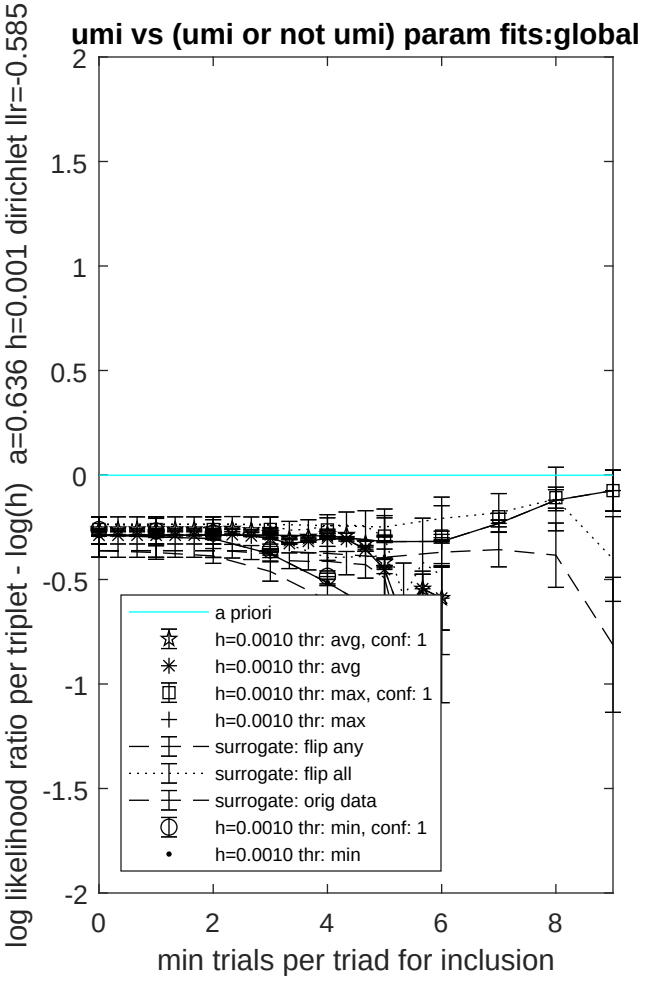

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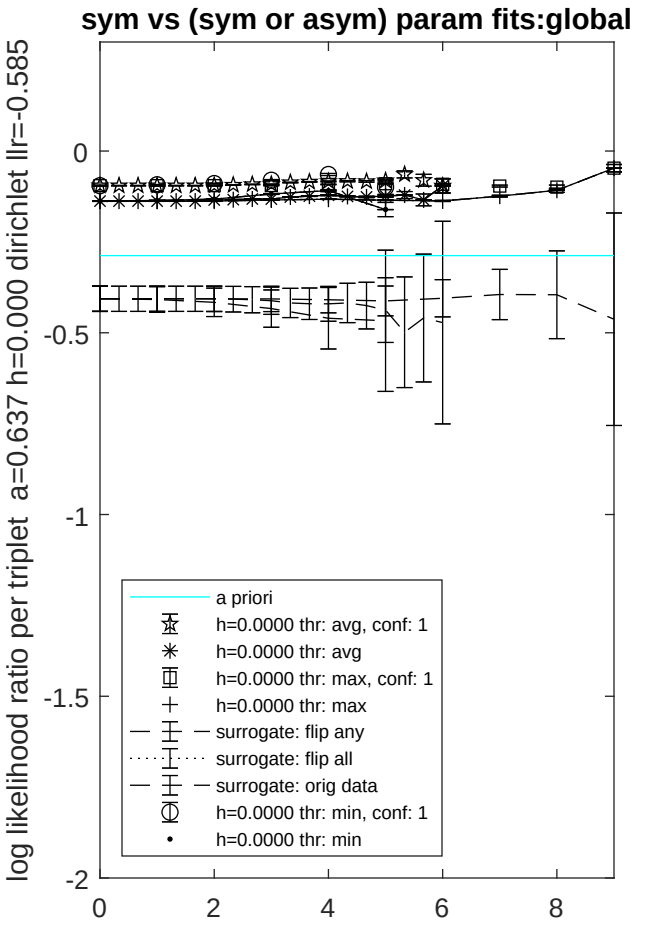

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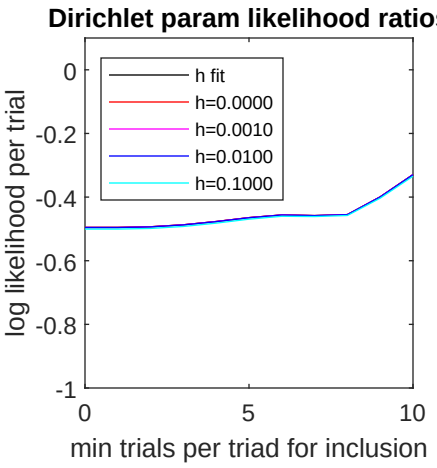

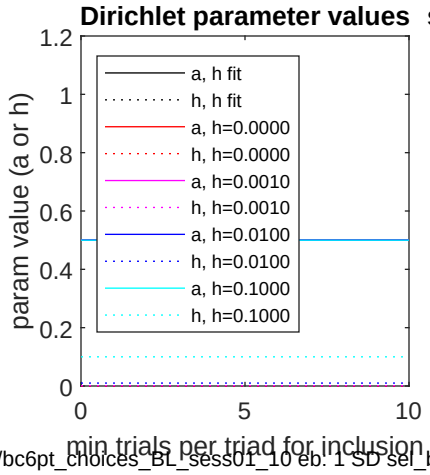

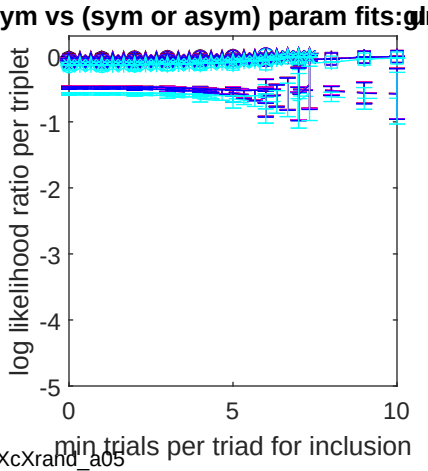

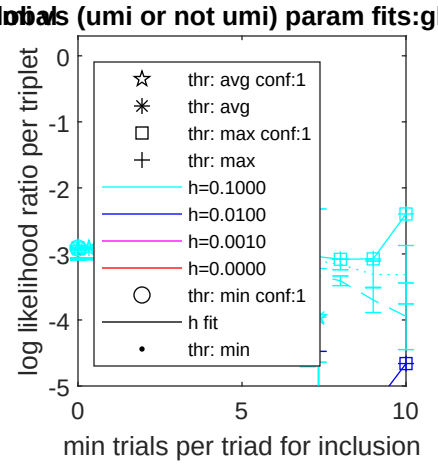

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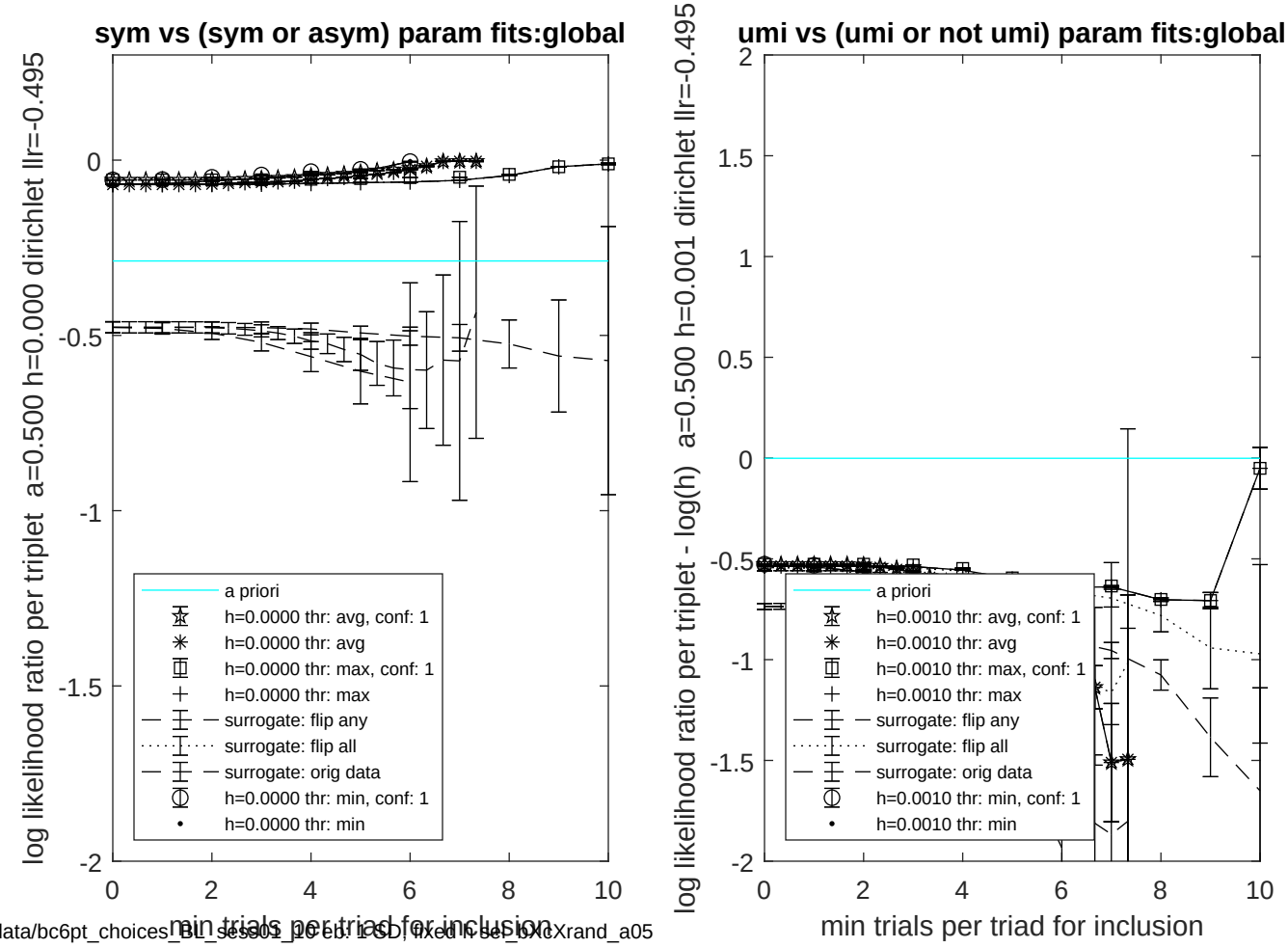

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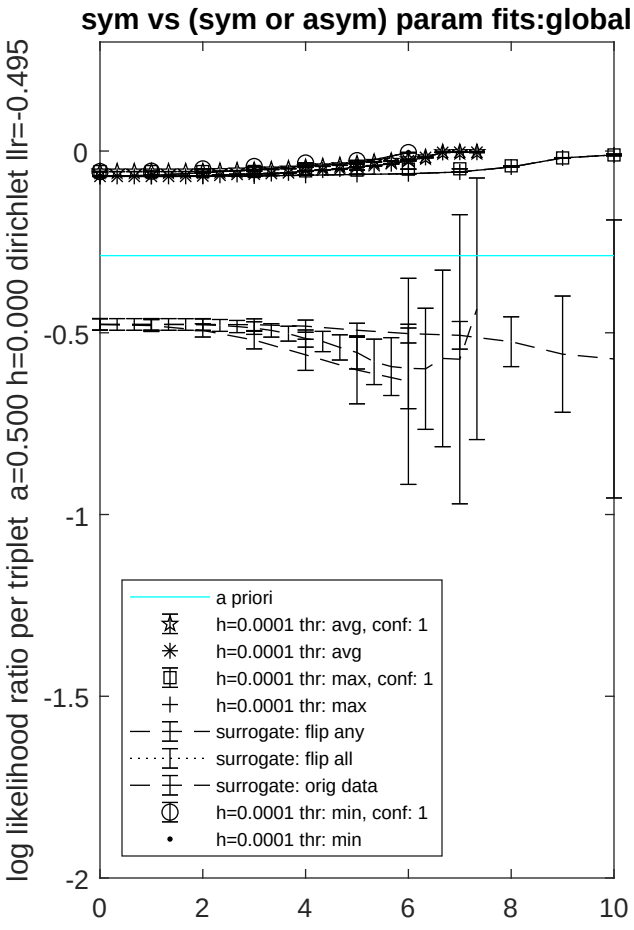
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-1
-1.5
-2
0
2
4
6
8
10
min trials per triad for inclusion
a priori
h=0.0001 thr: avg, conf: 1
h=0.0001 thr: avg
h=0.0001 thr: max, conf: 1
h=0.0001 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0001 thr: min, conf: 1
h=0.0001 thr: min
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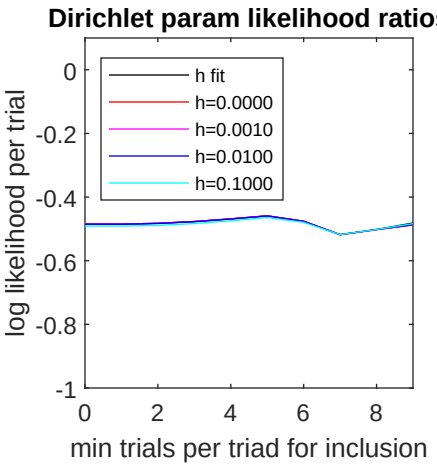

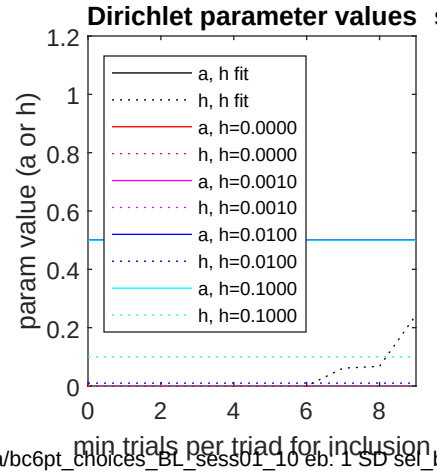

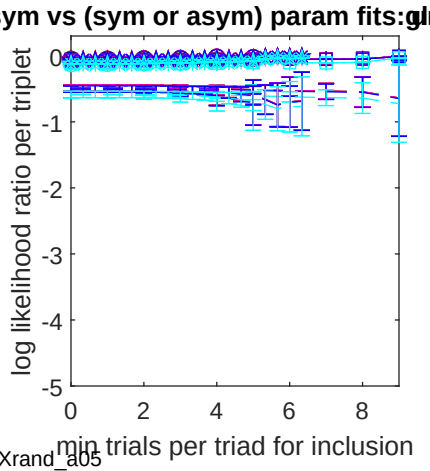

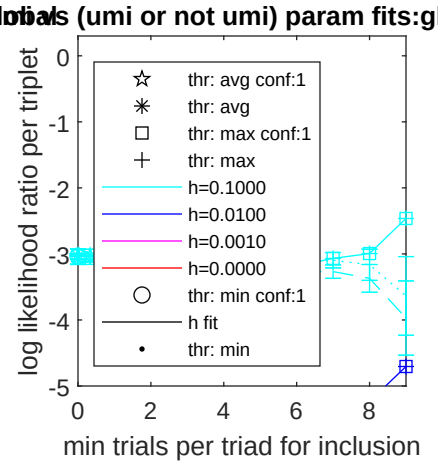

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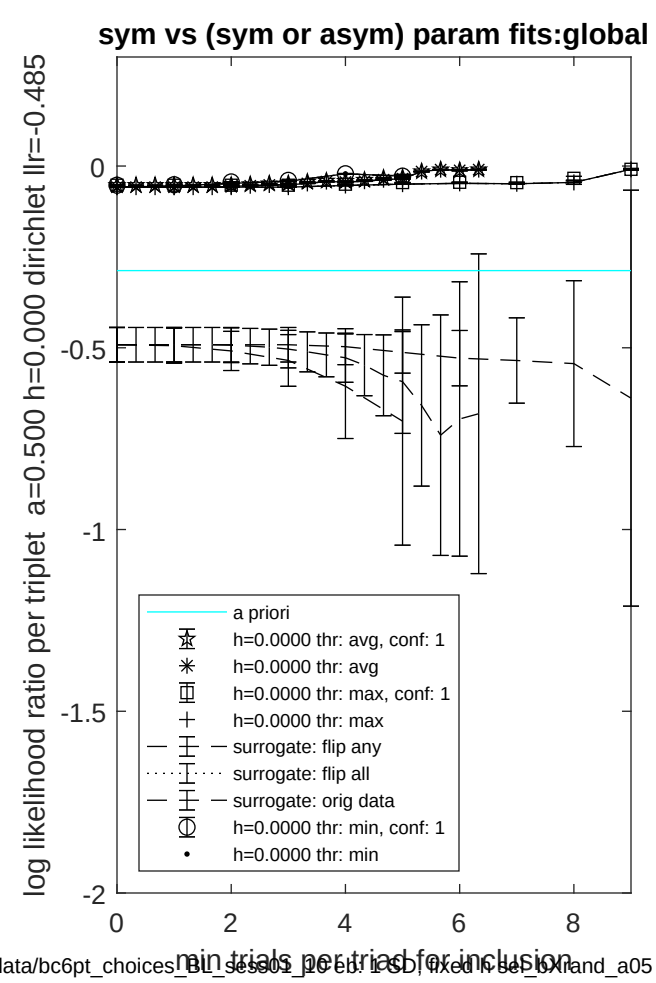

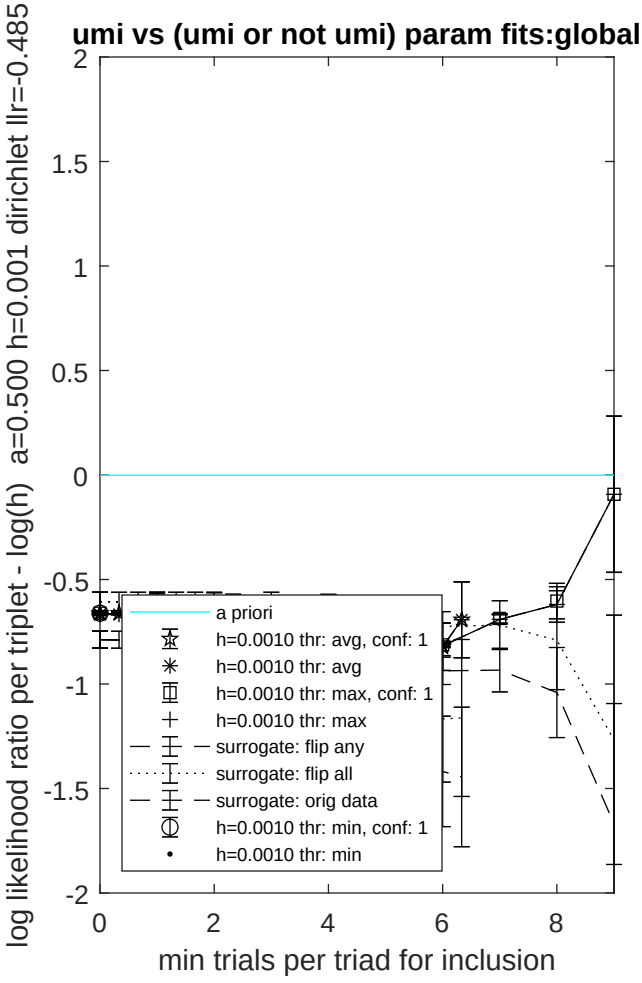

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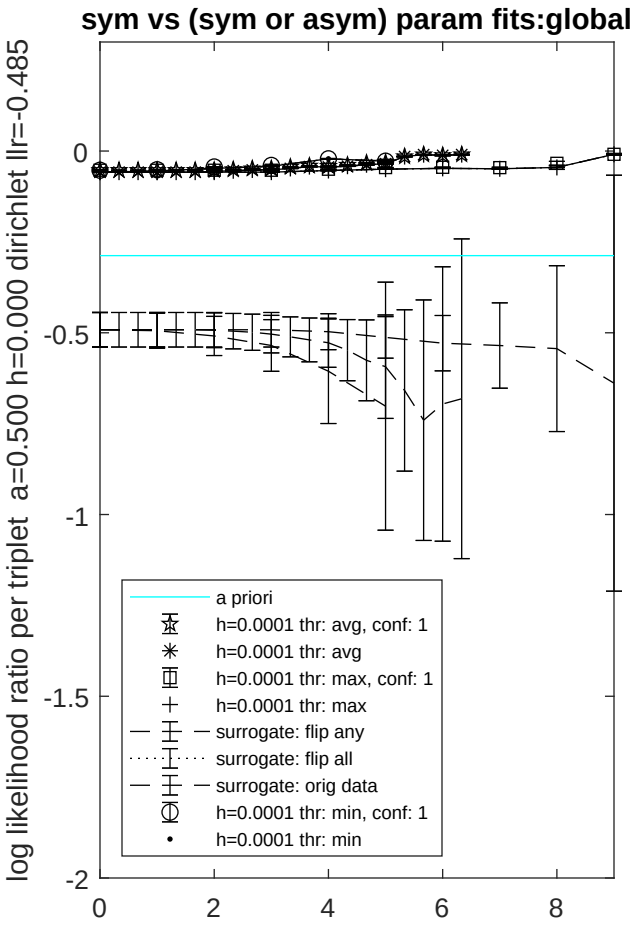

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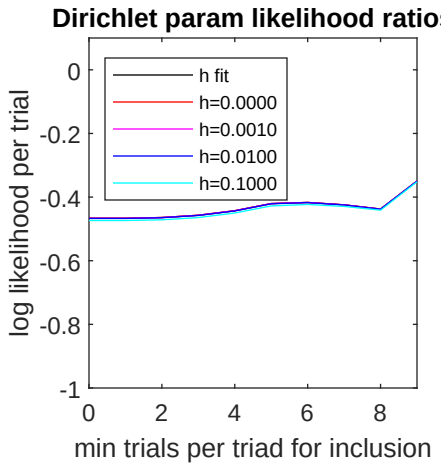

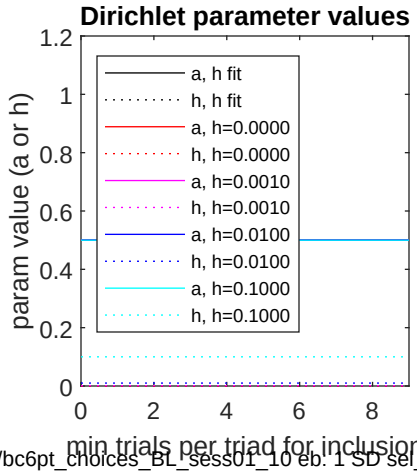

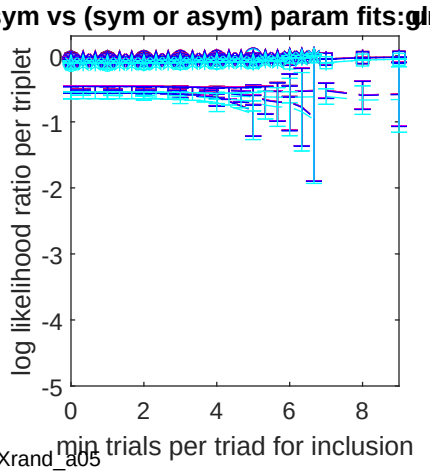

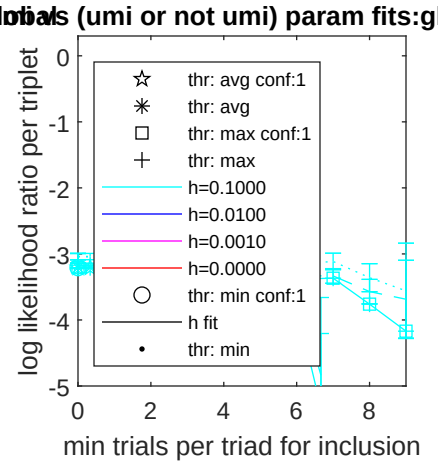

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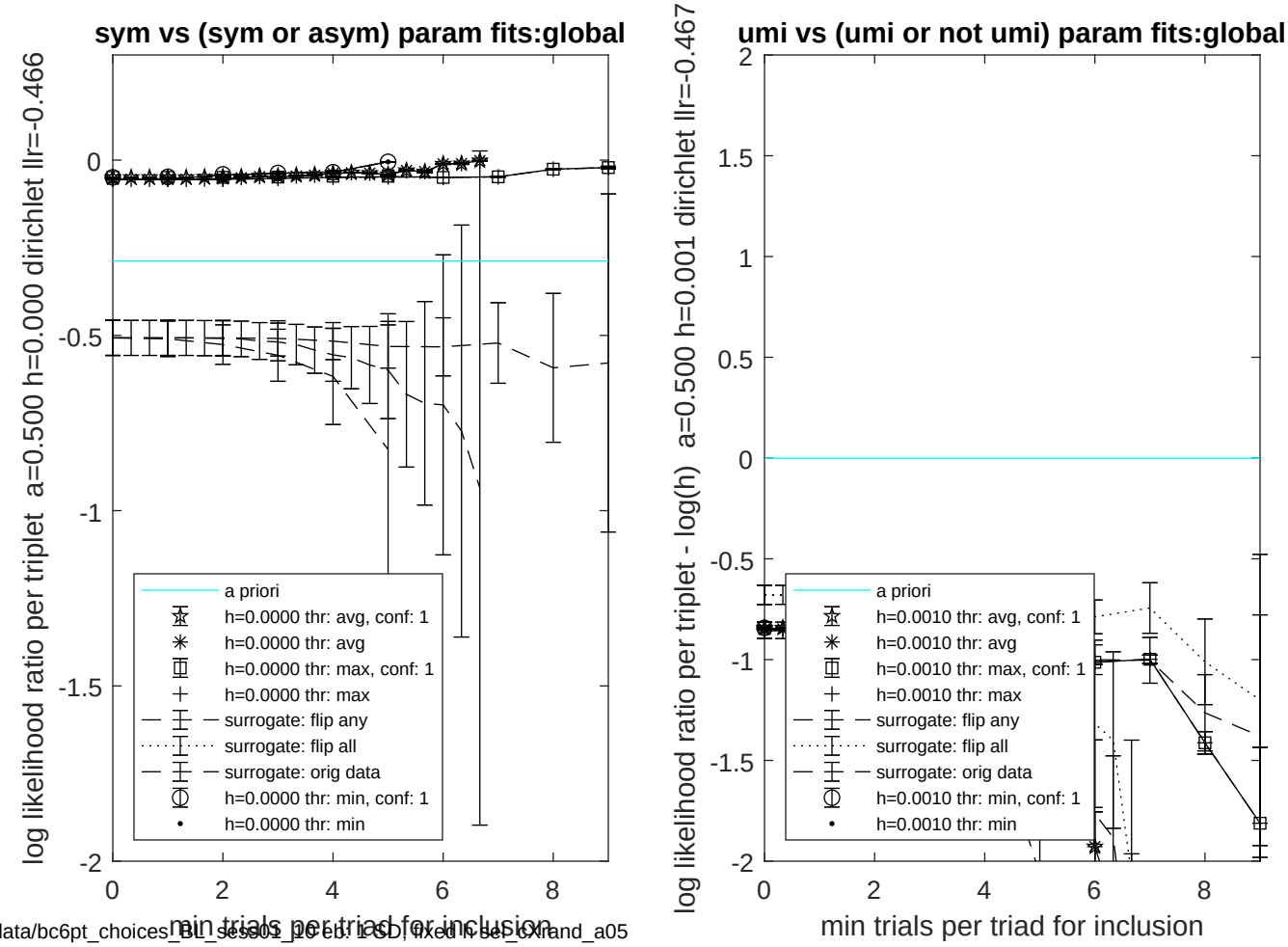

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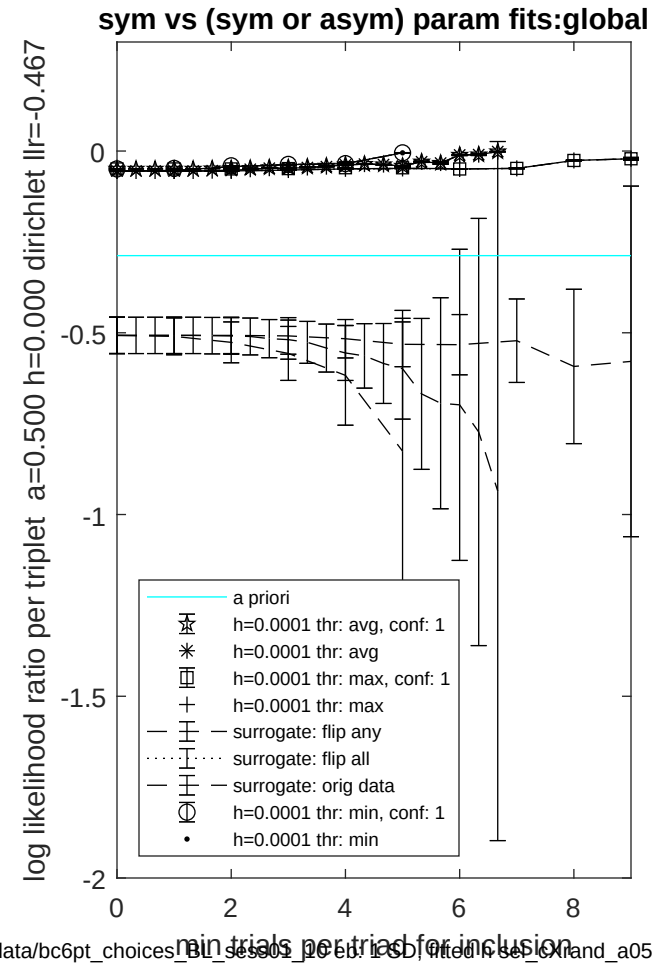

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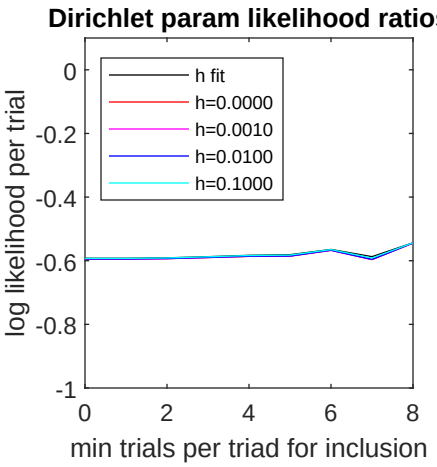

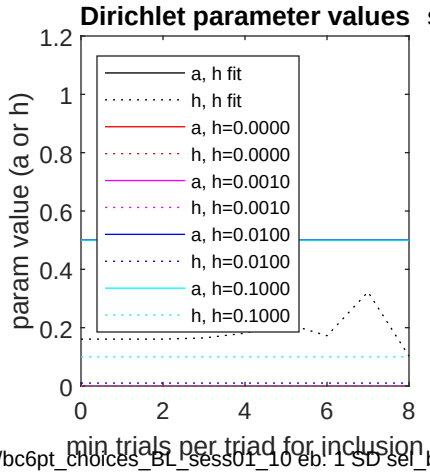

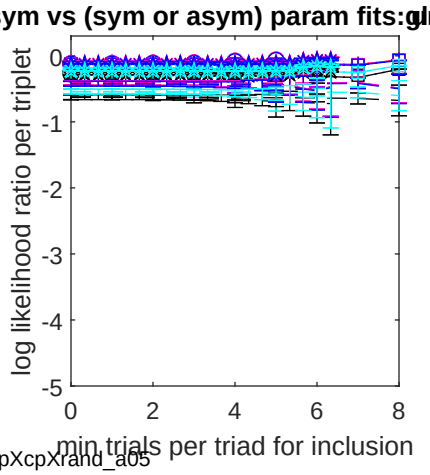

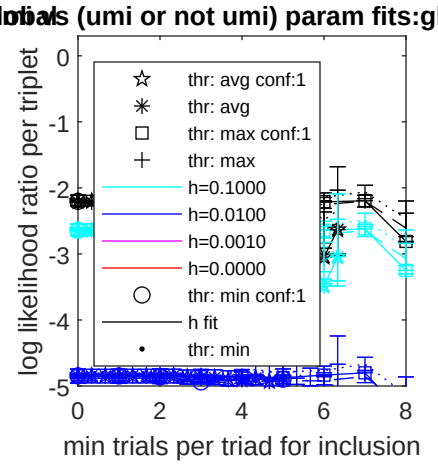

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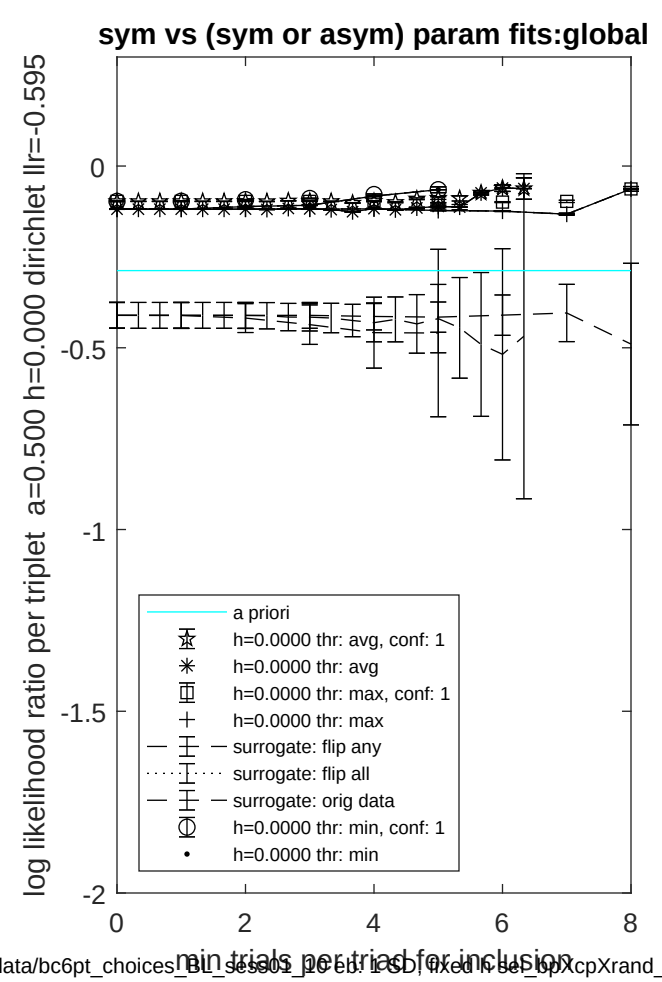

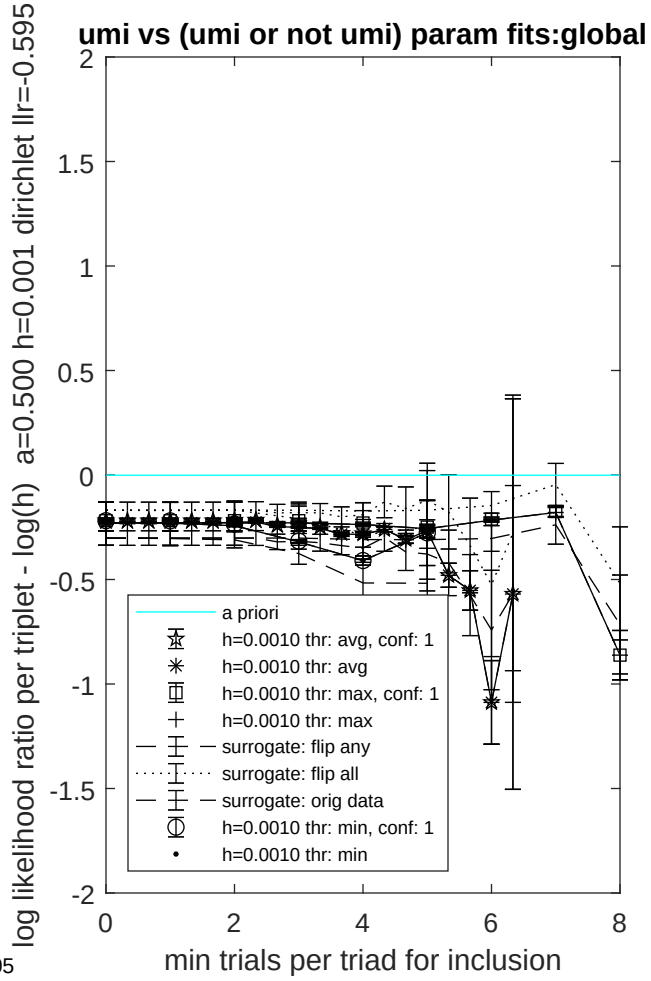

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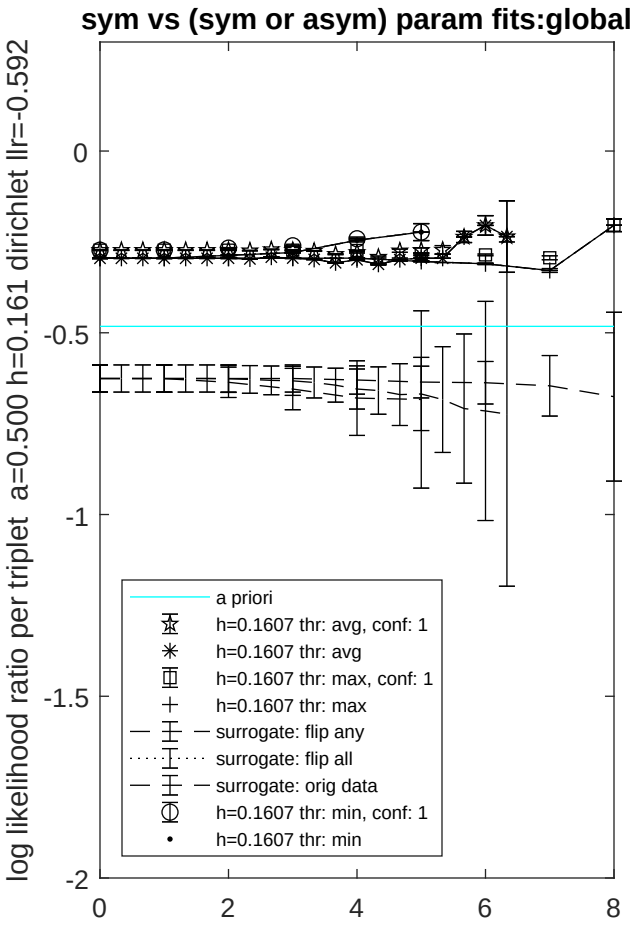

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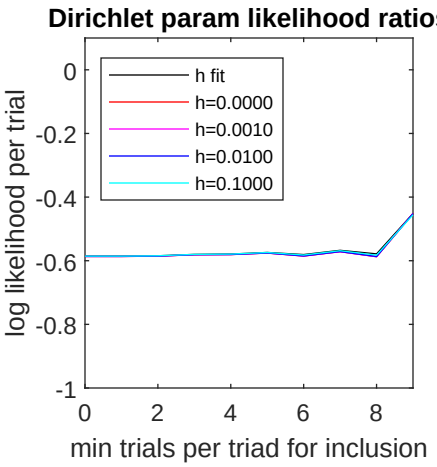

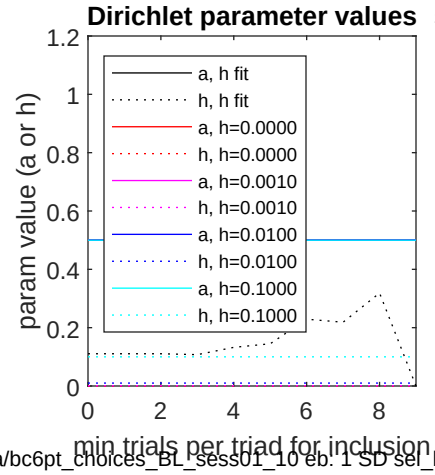

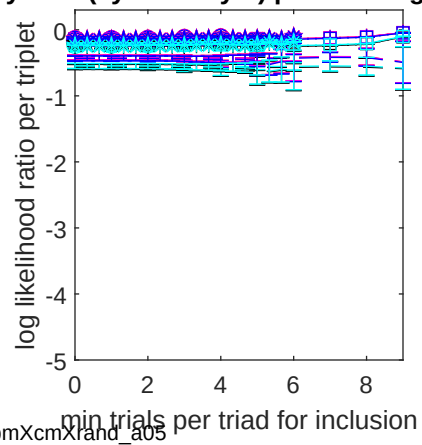

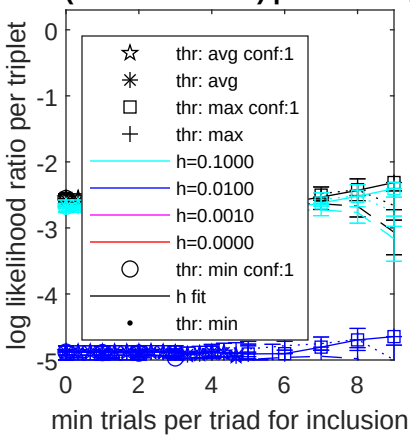

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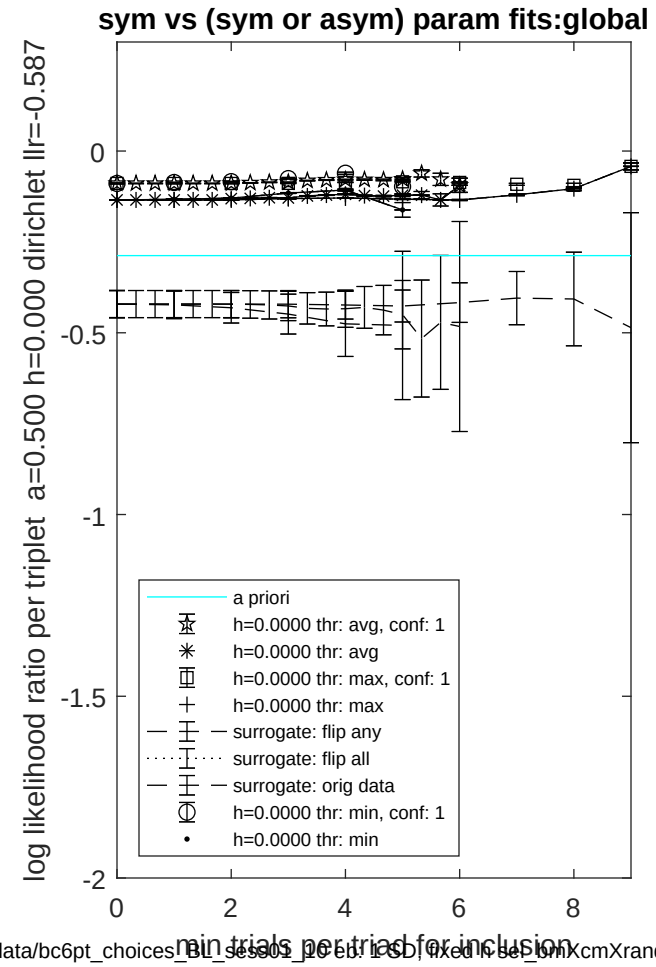

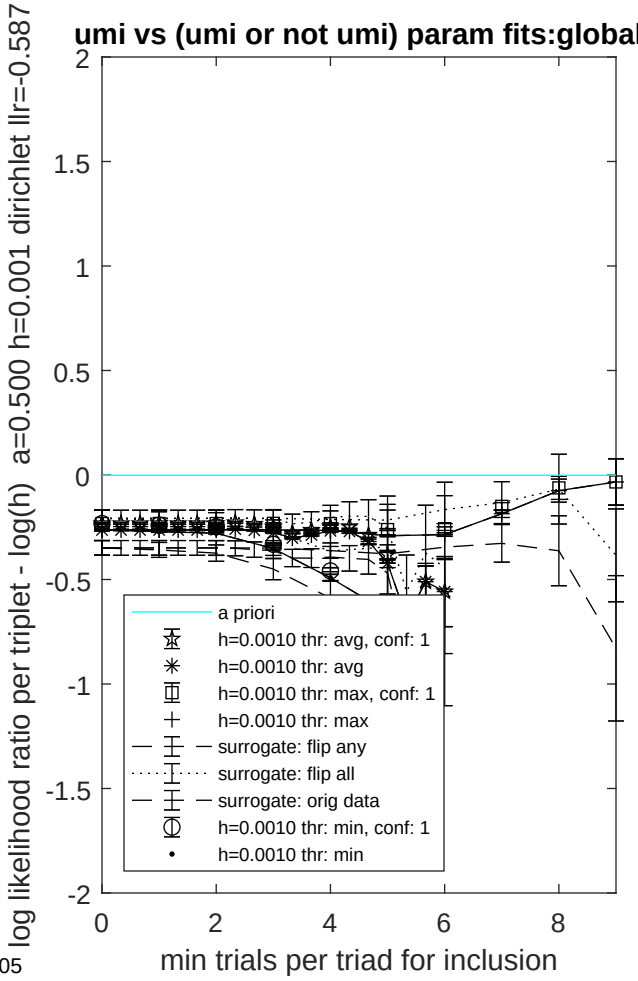

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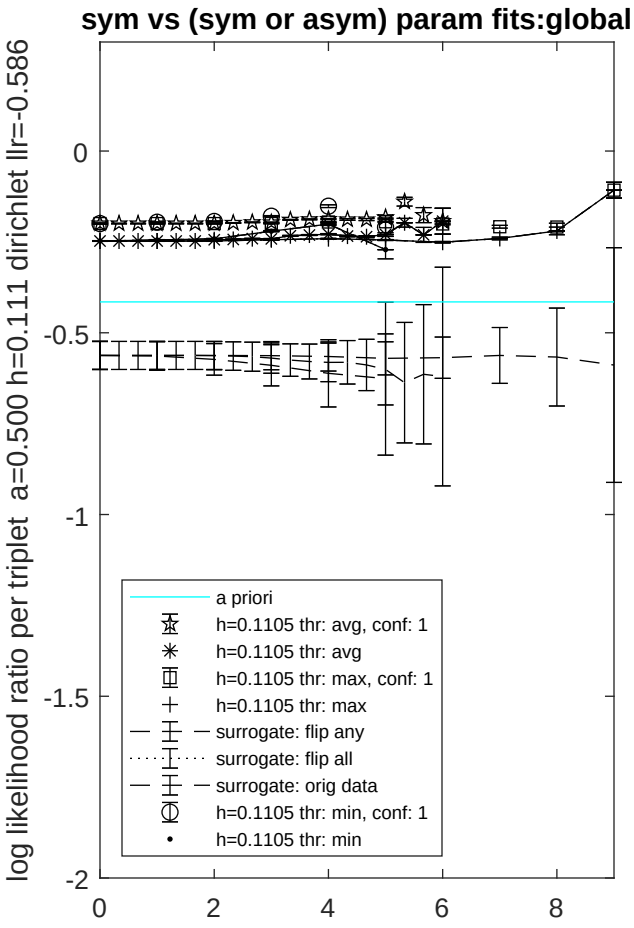

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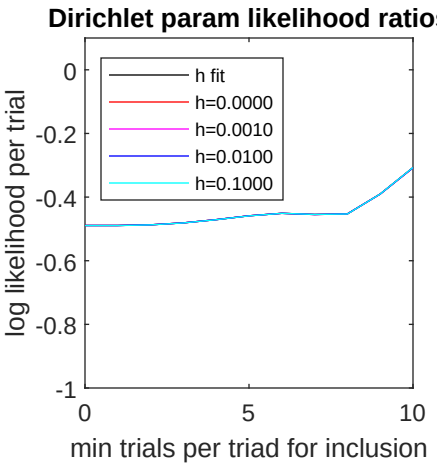

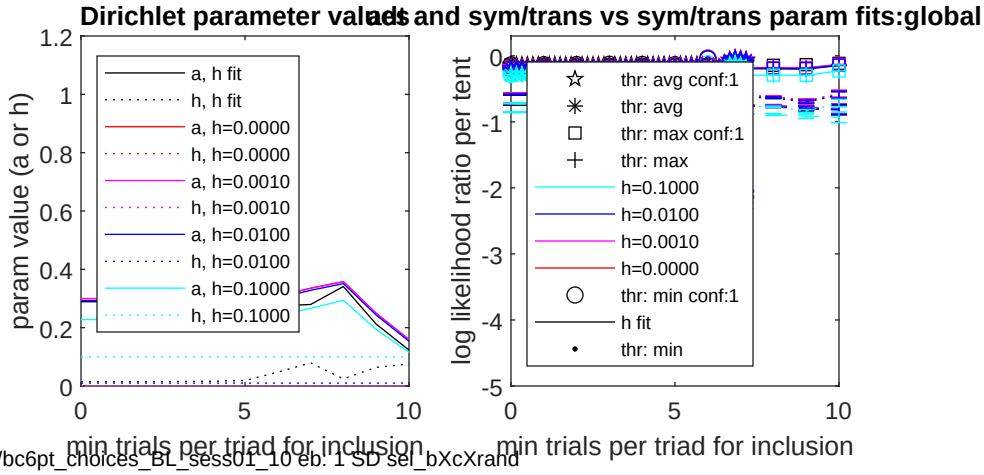

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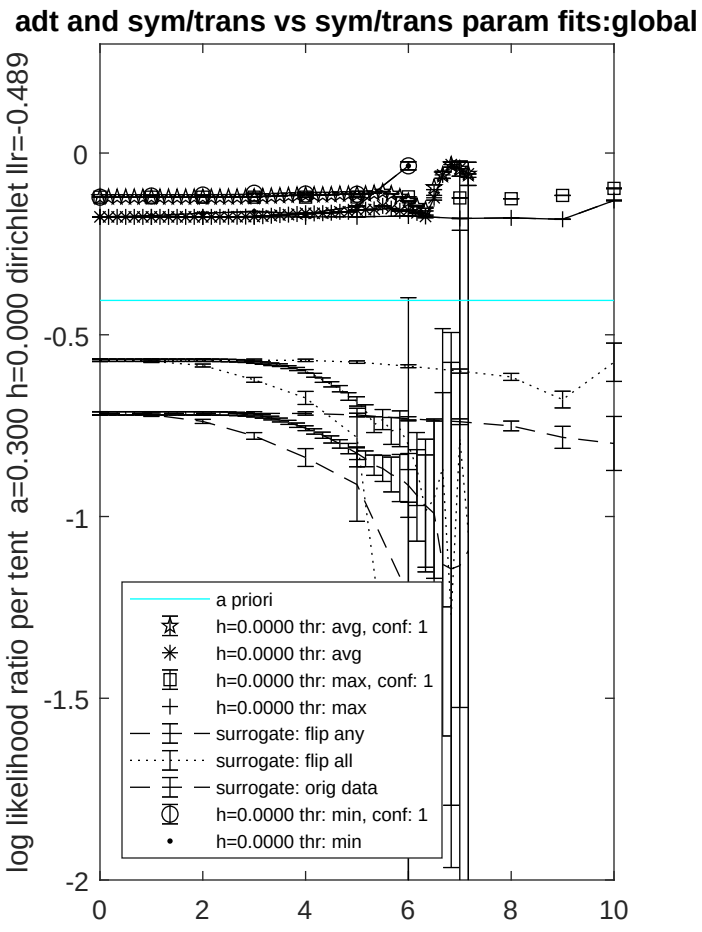

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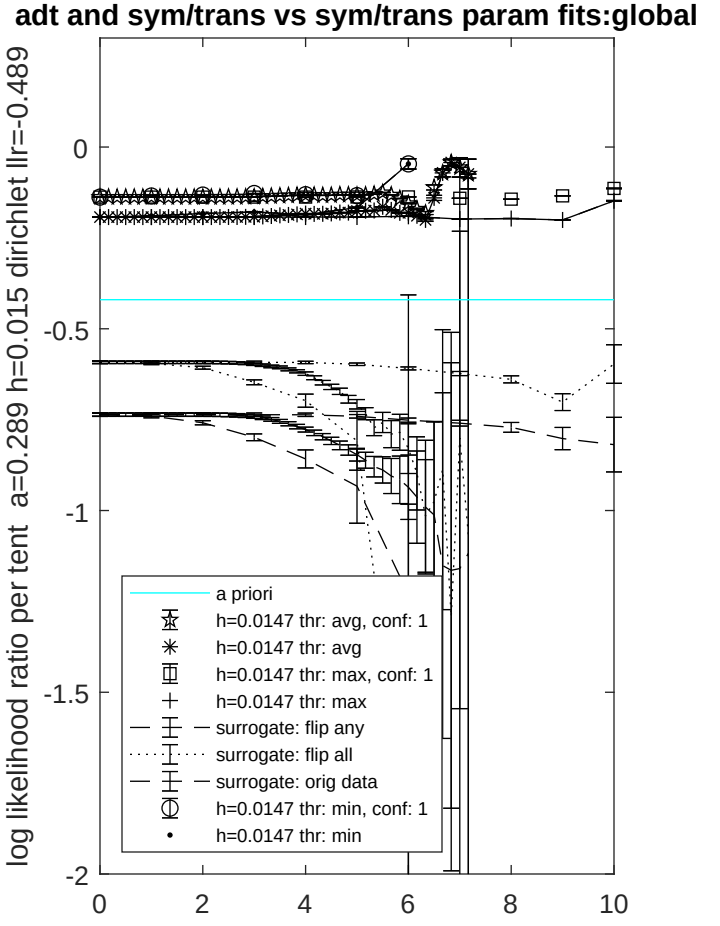

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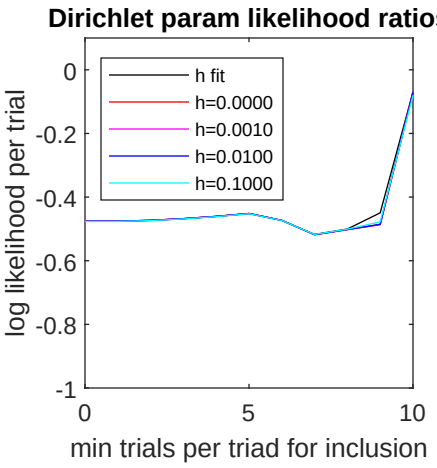

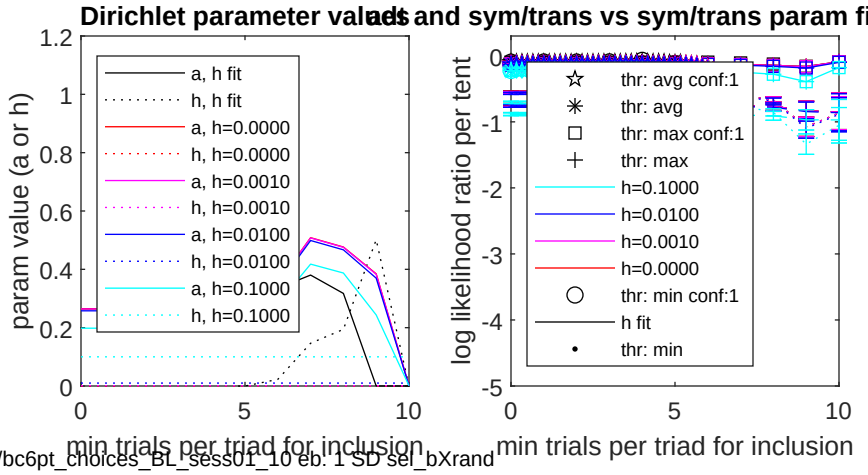

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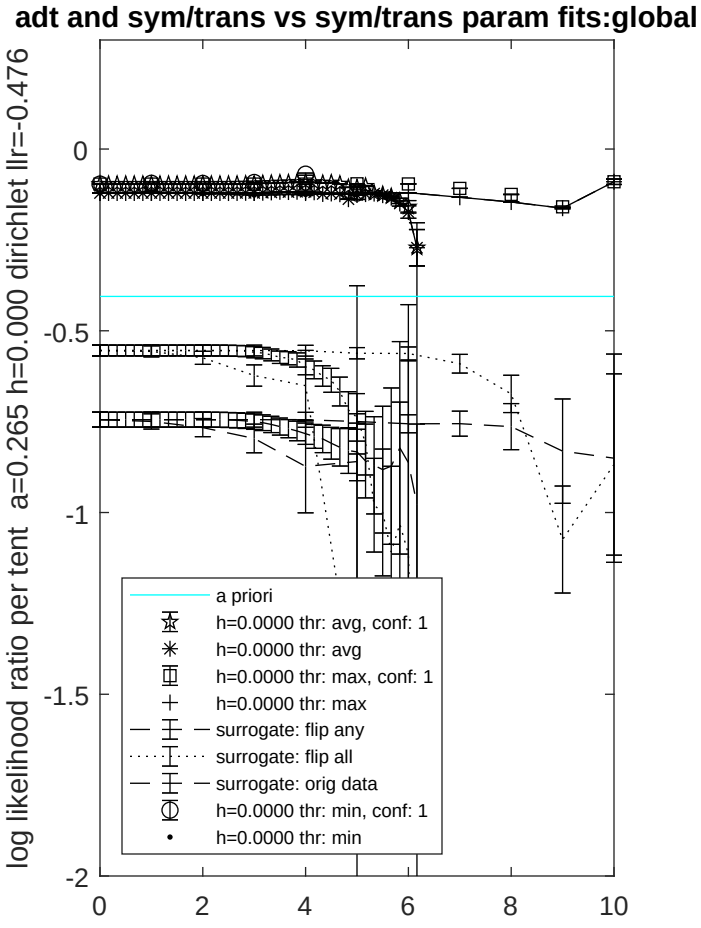

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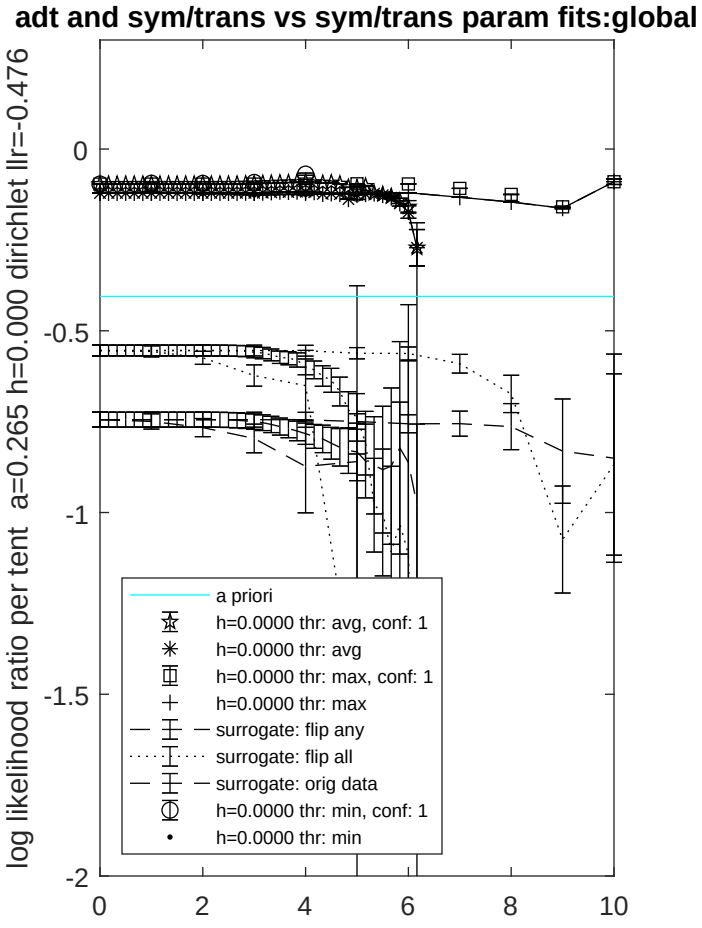

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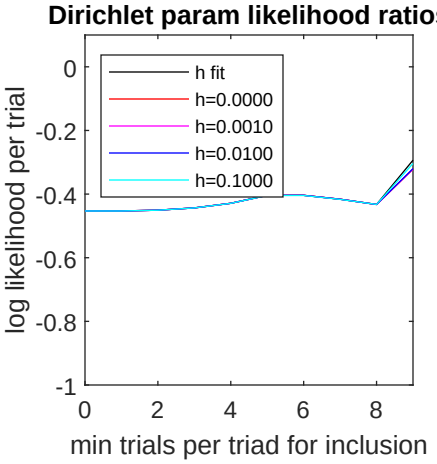

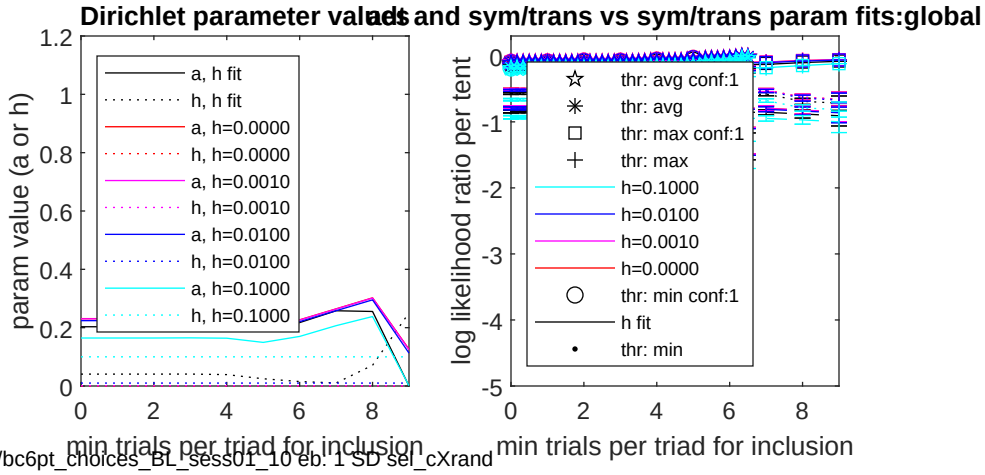

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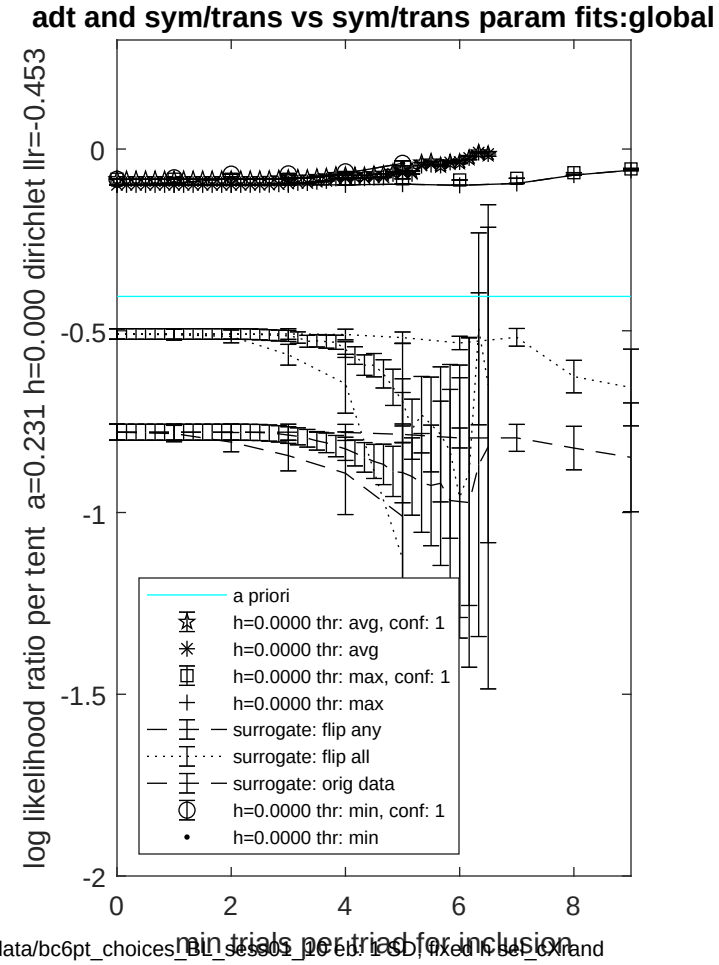

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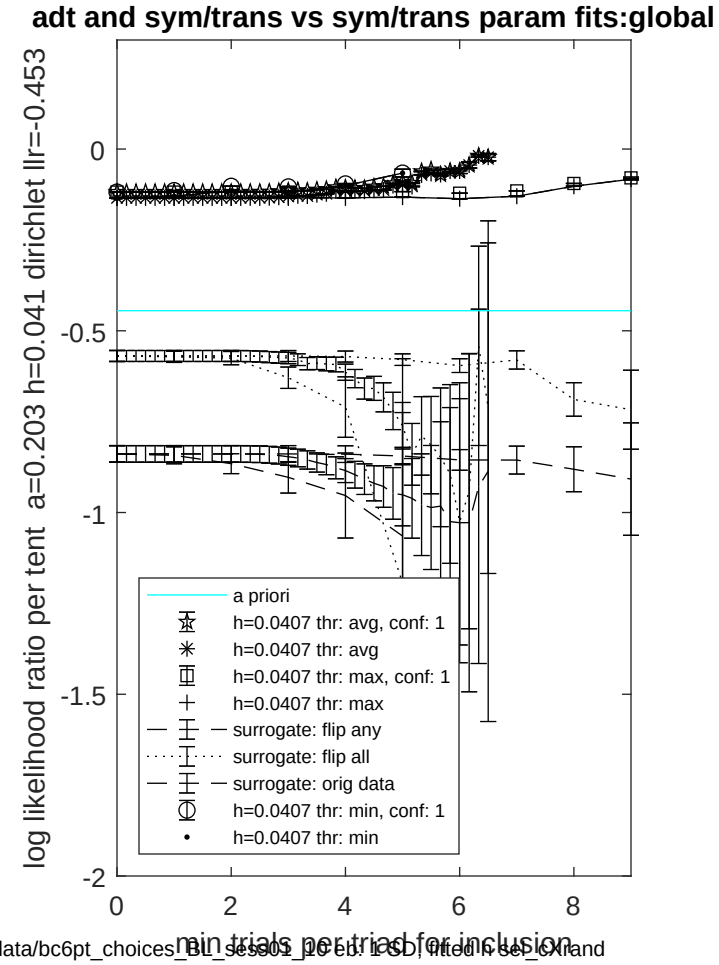

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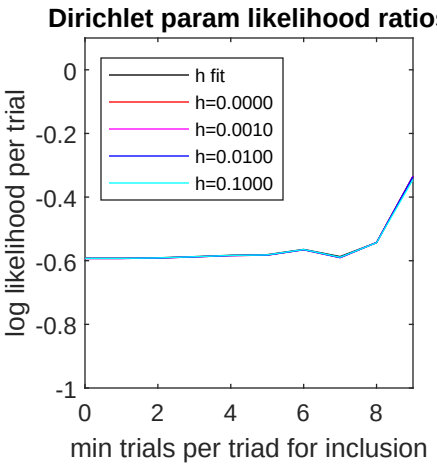

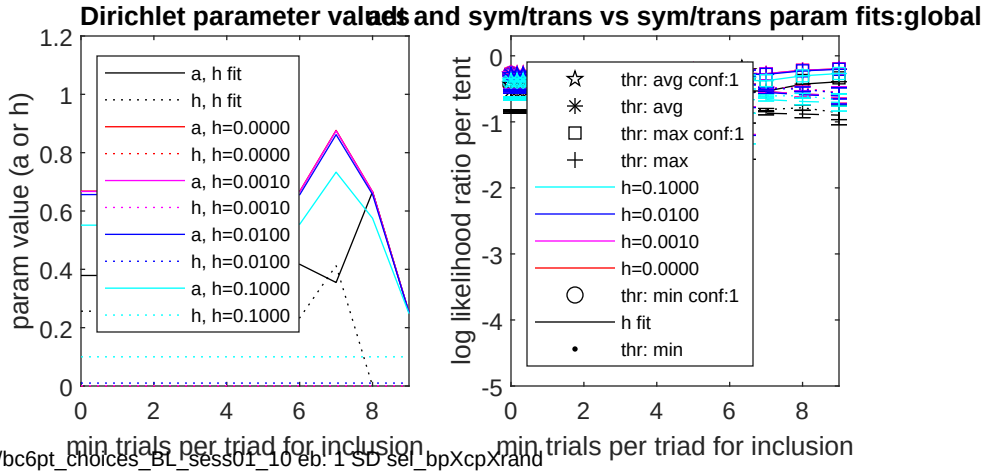

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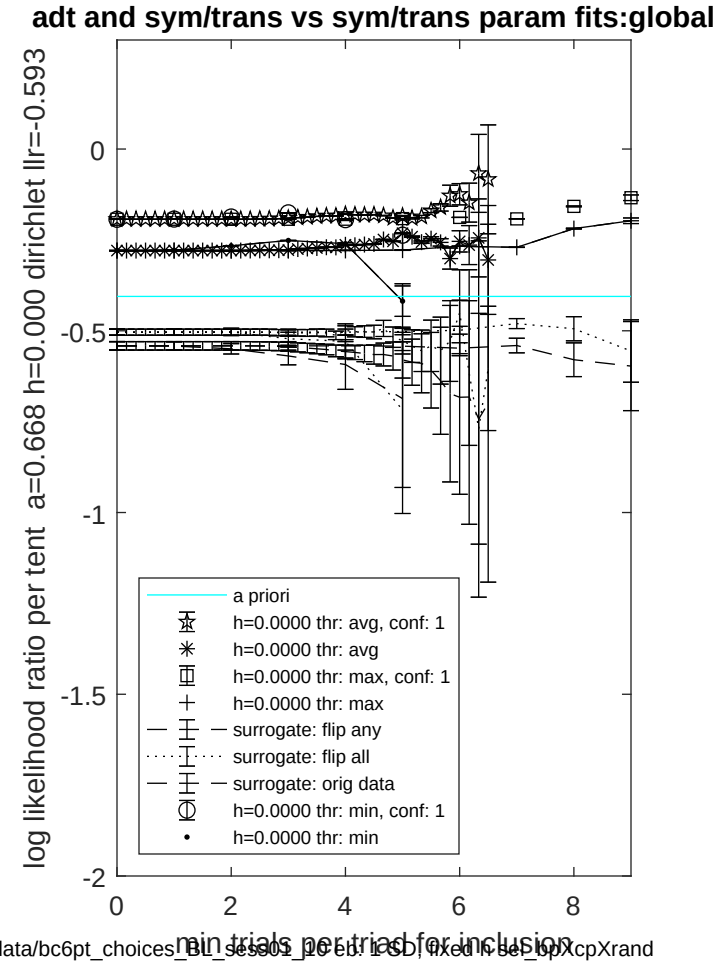

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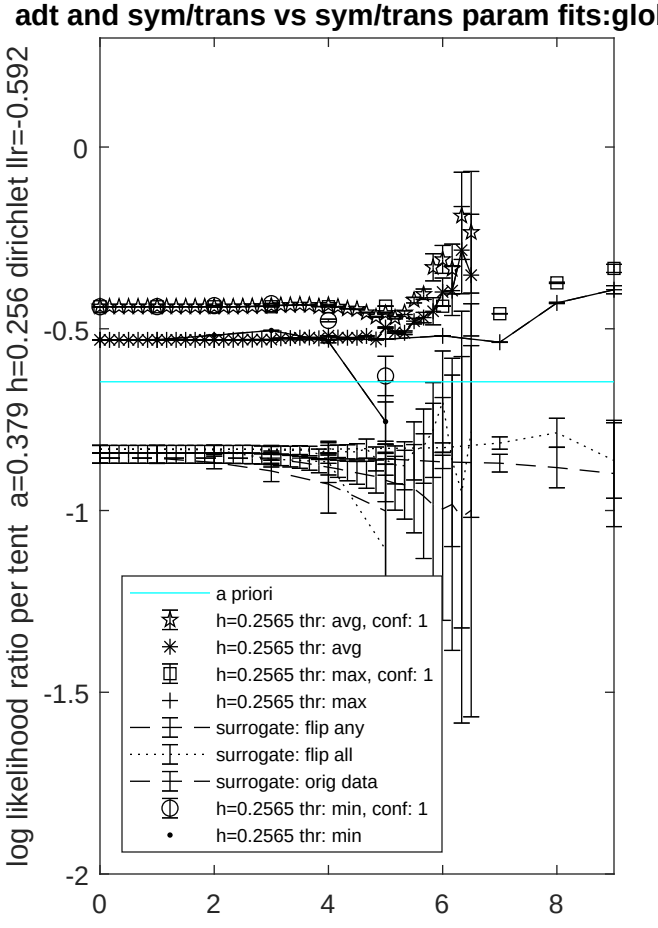

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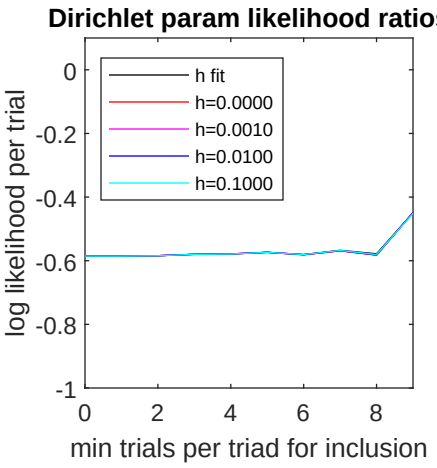

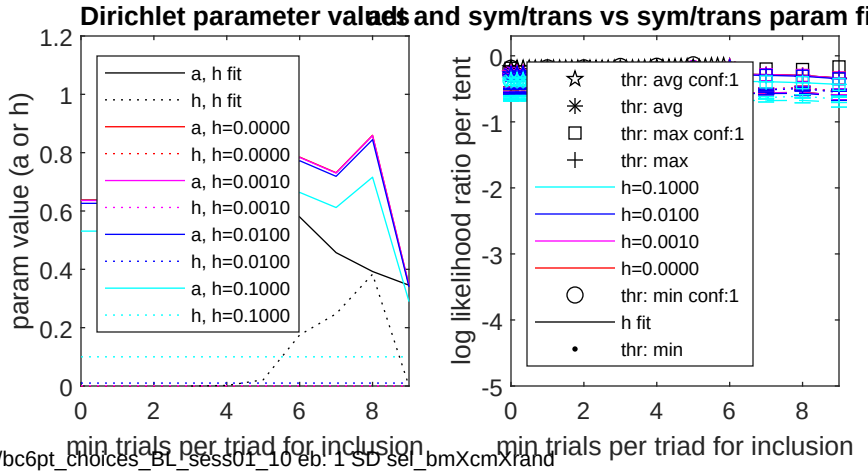

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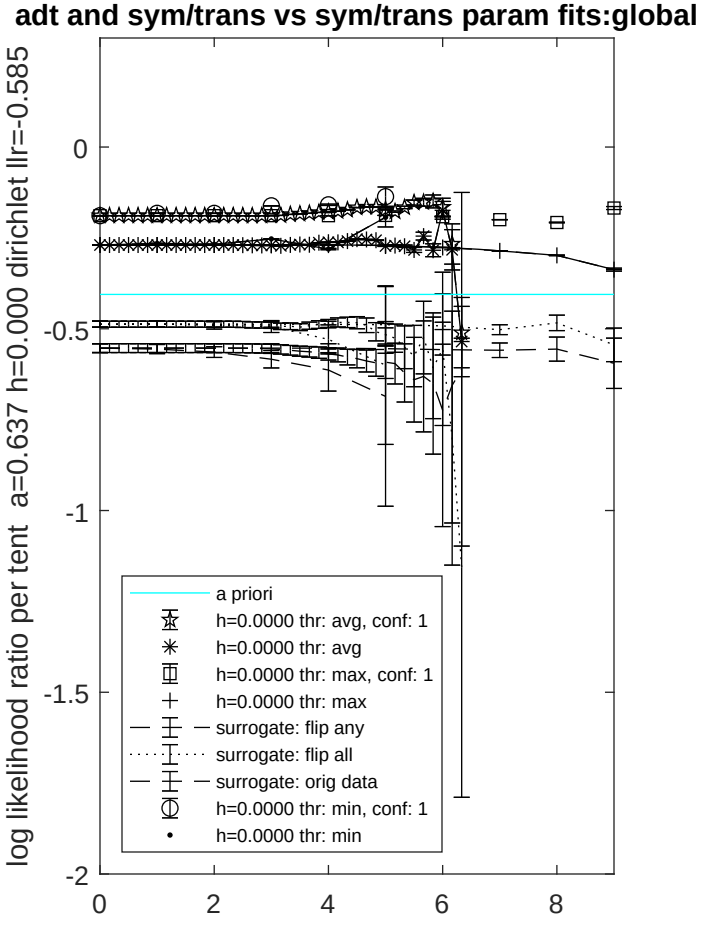

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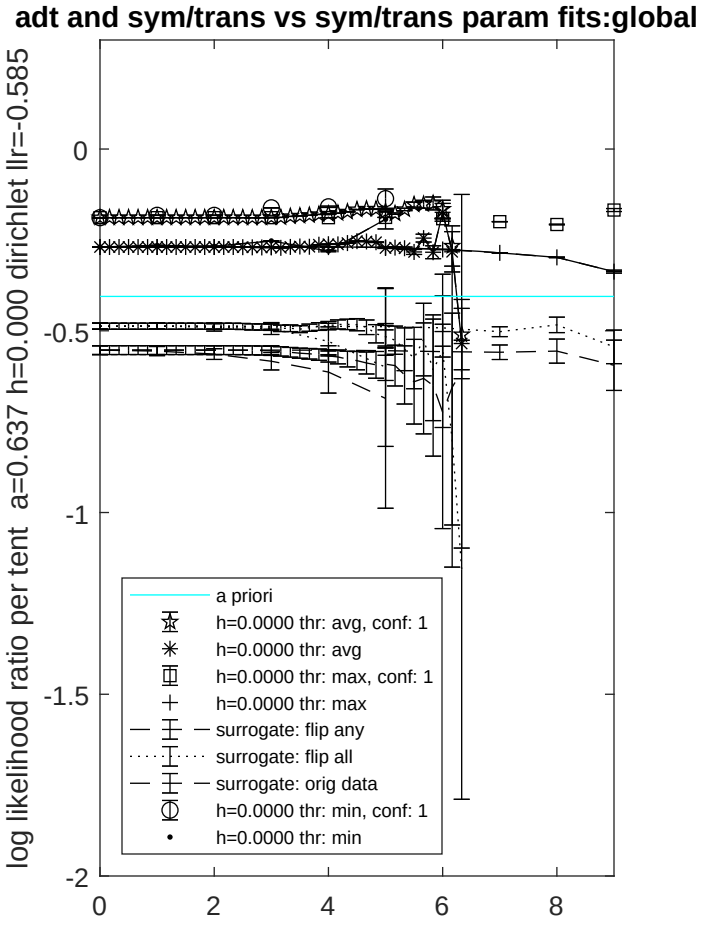

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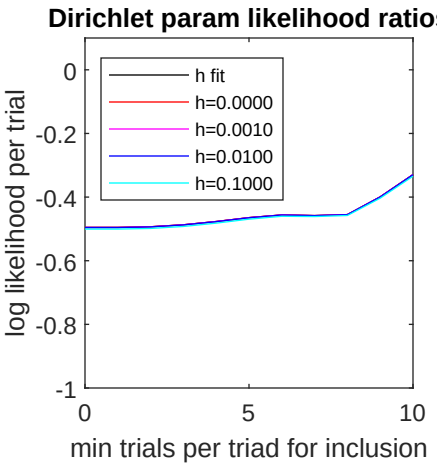

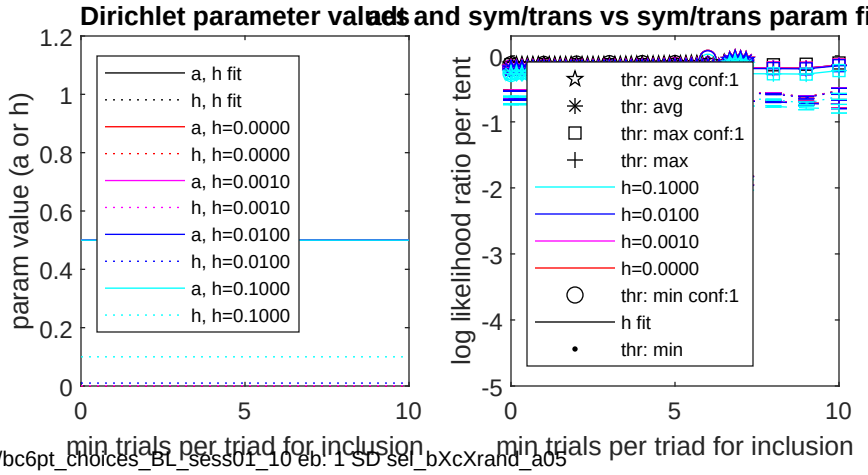

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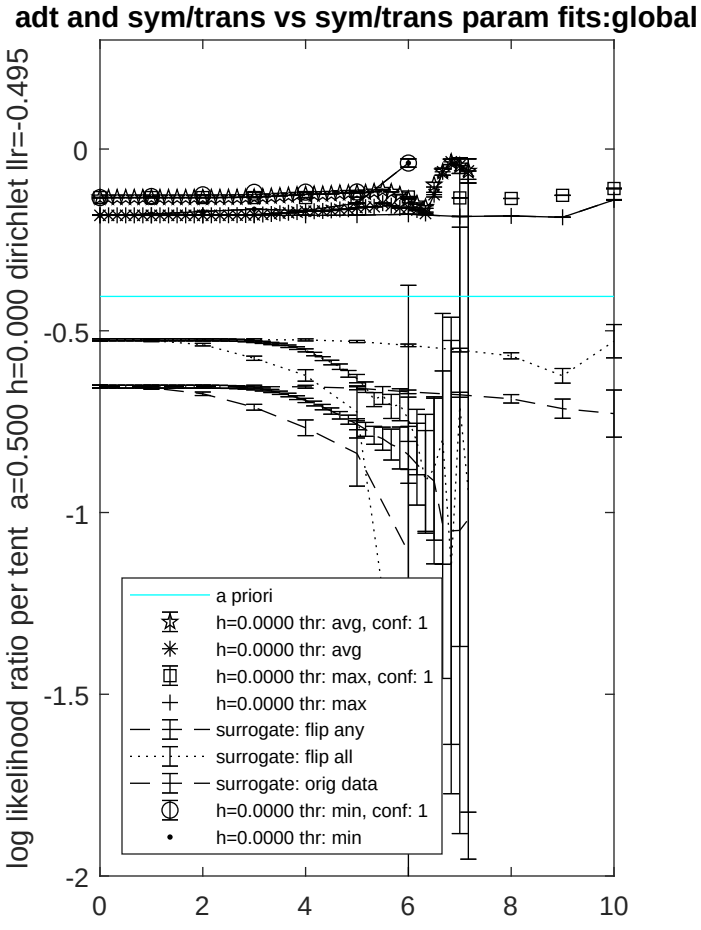

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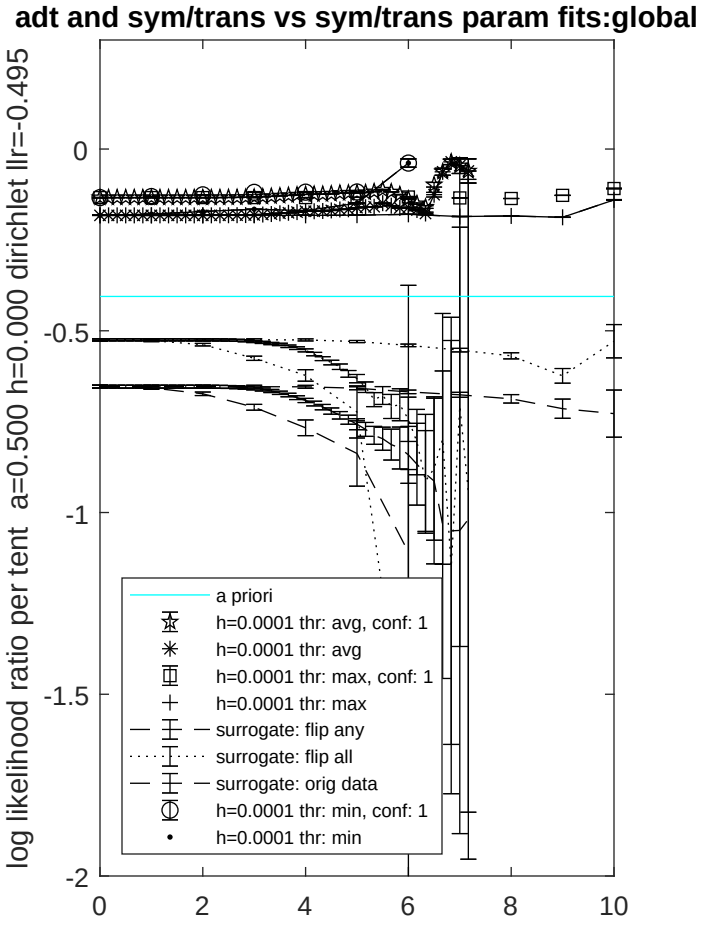
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-0.5
-1
-1.5
-2
0
2
4
6
8
10
min trials per triad for inclusion
a priori
h=0.0001 thr: avg, conf: 1
h=0.0001 thr: avg
h=0.0001 thr: max, conf: 1
h=0.0001 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0001 thr: min, conf: 1
h=0.0001 thr: min
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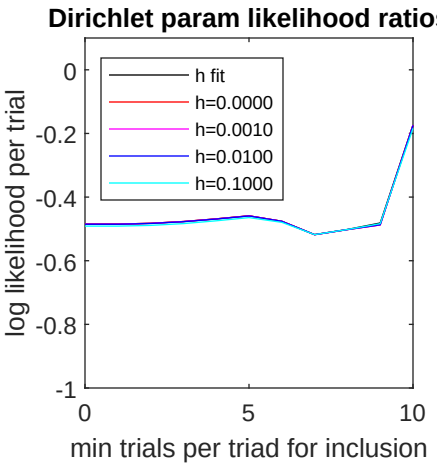

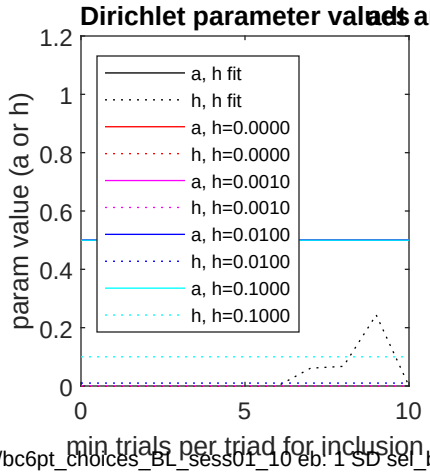

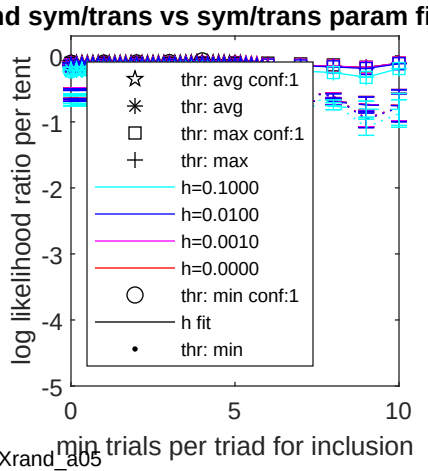

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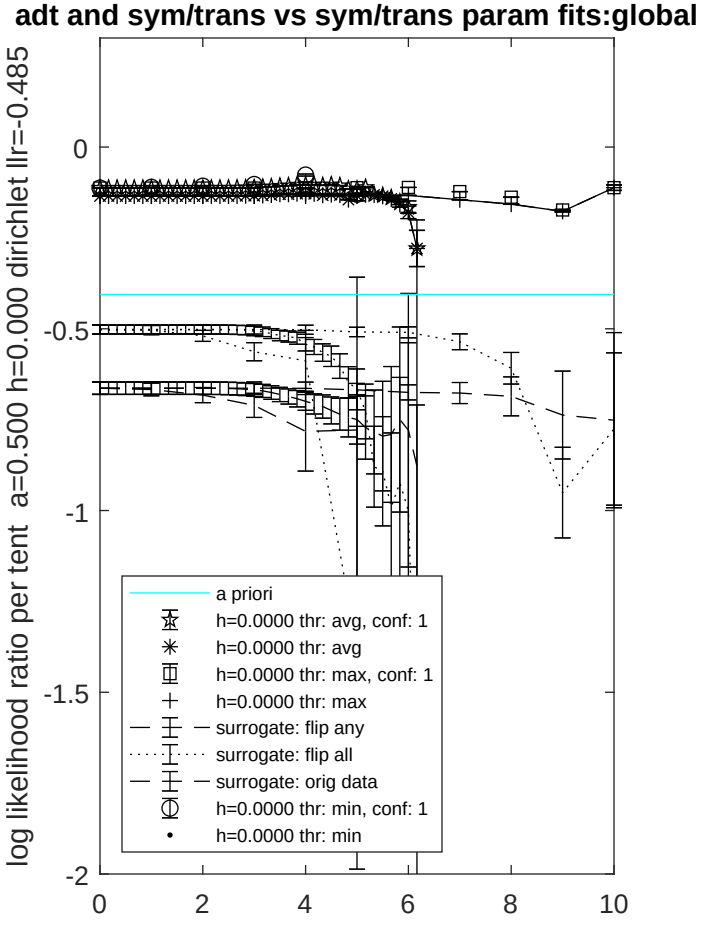

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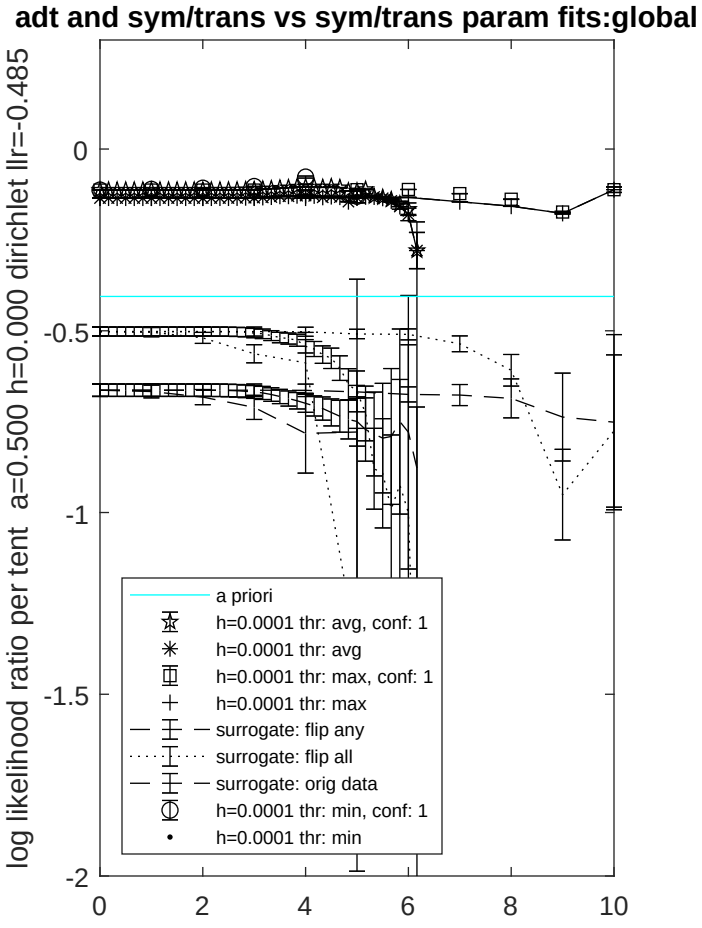

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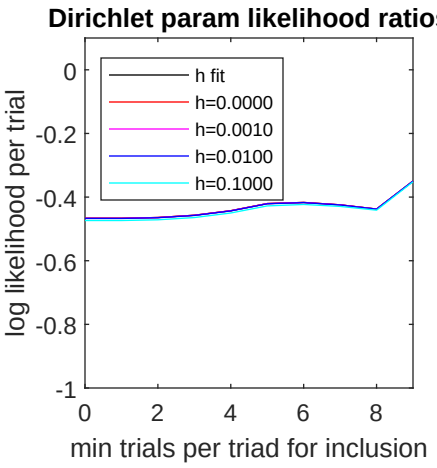

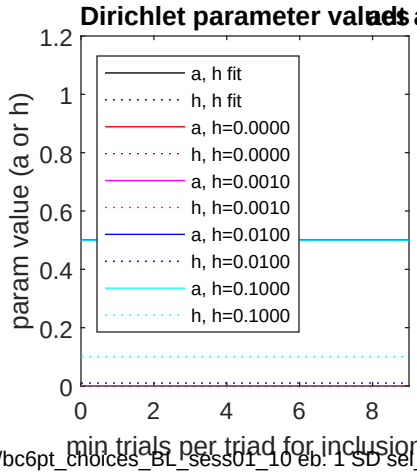

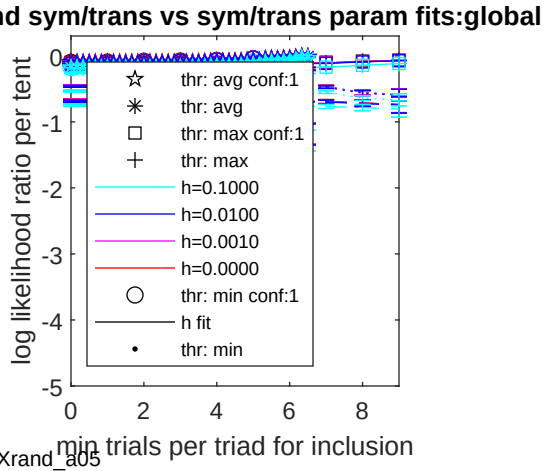

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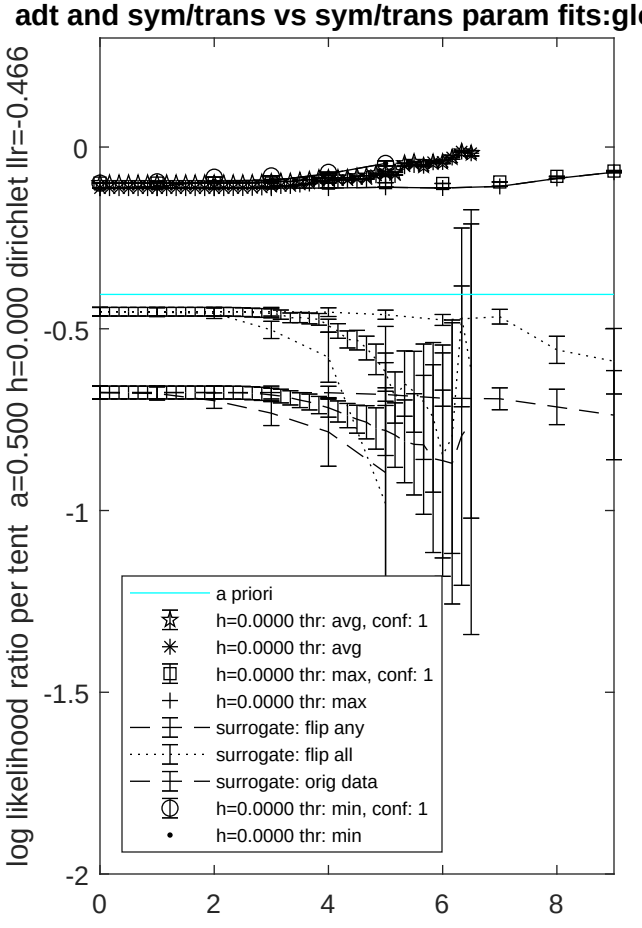

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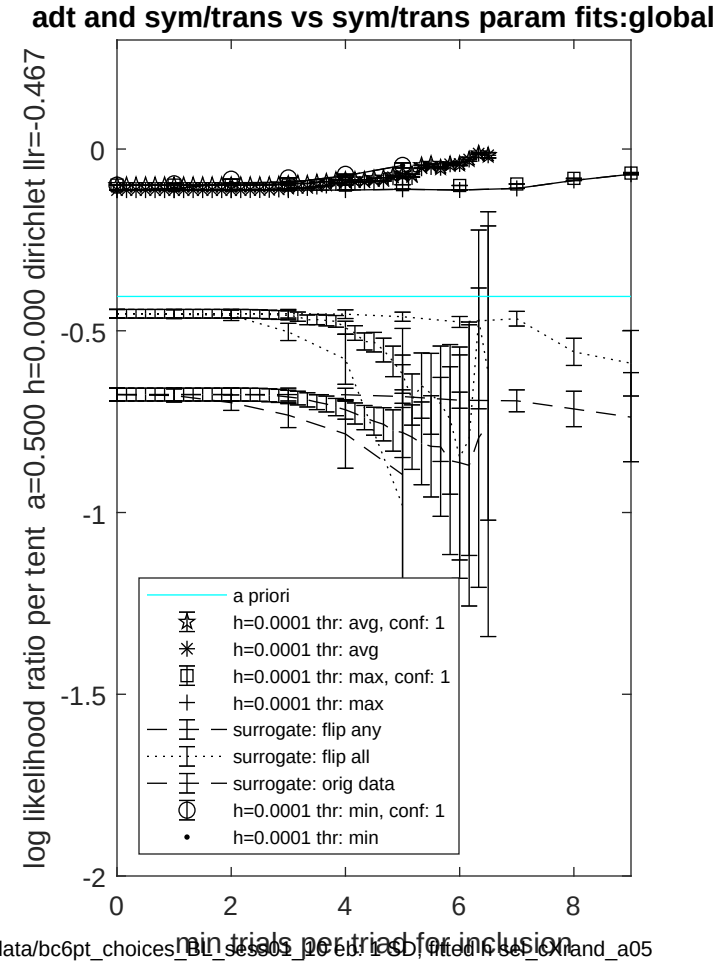

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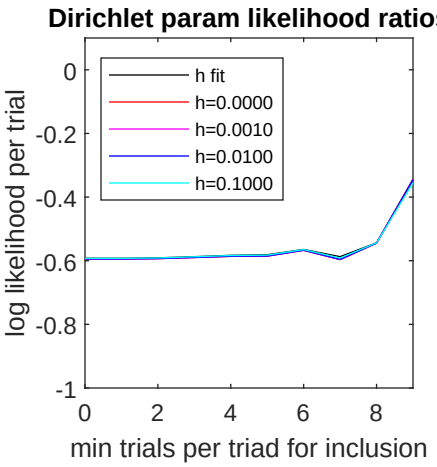

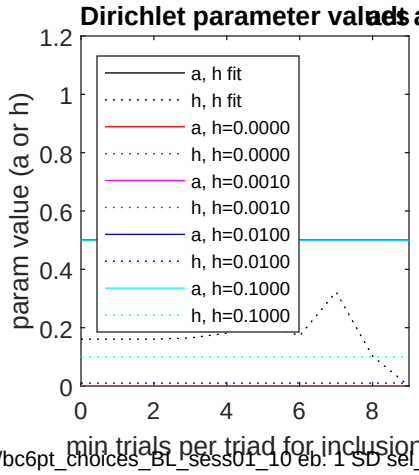

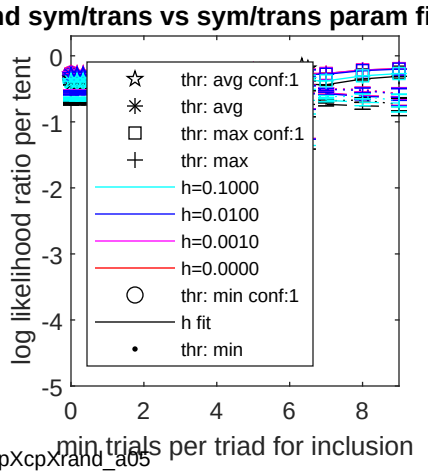

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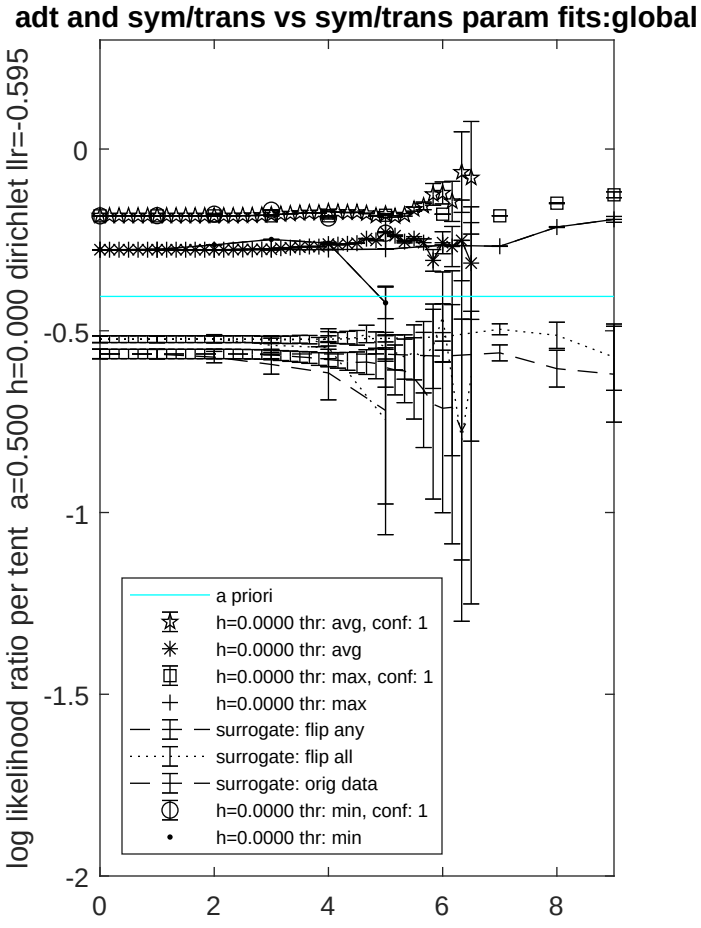

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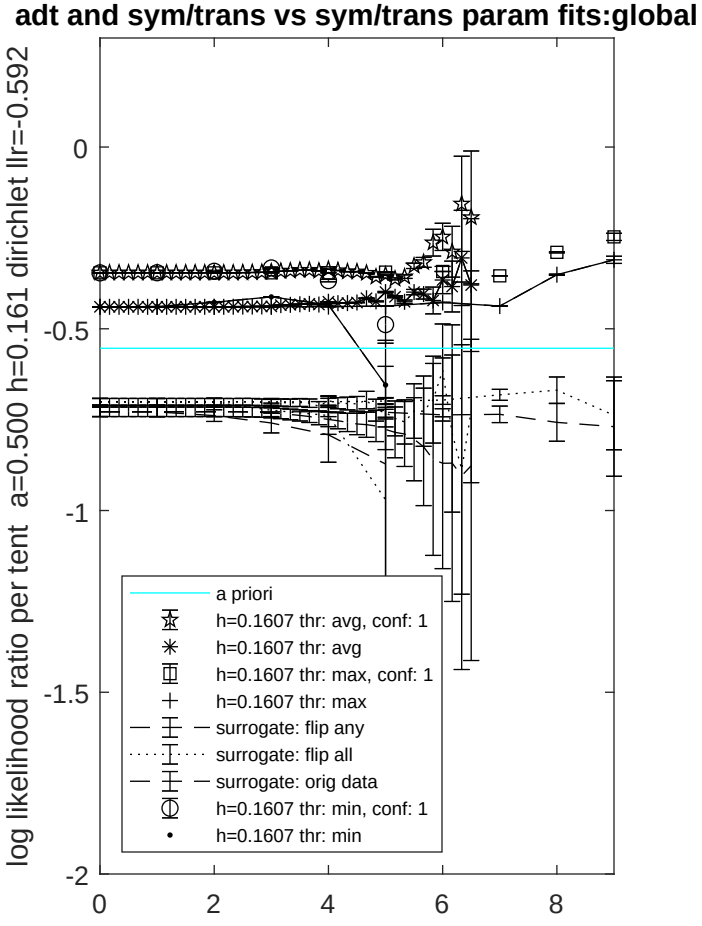

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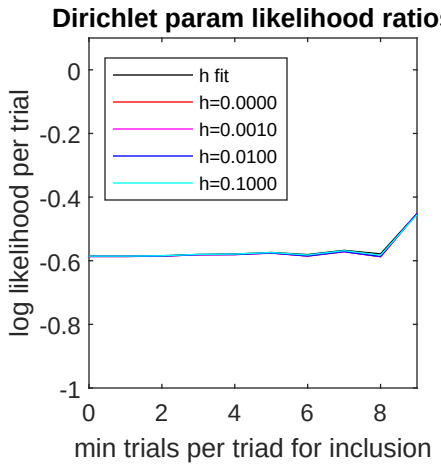

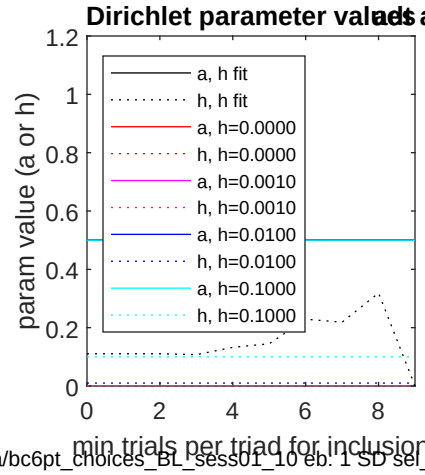

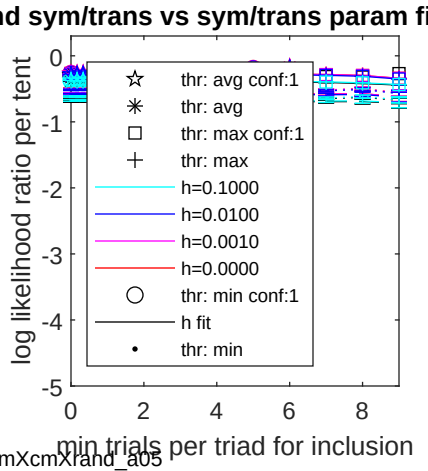

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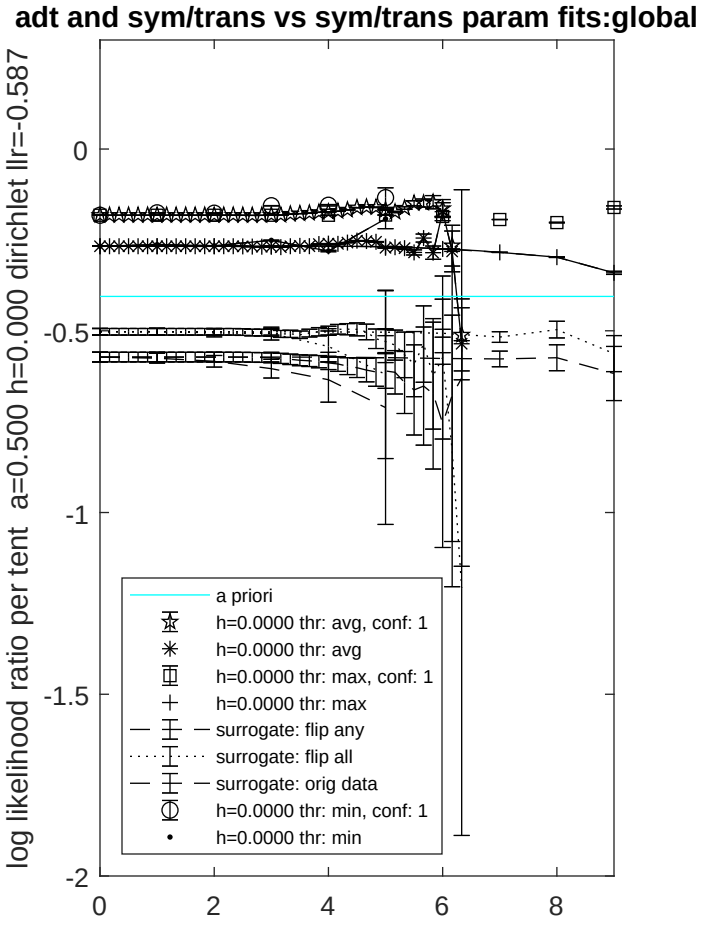

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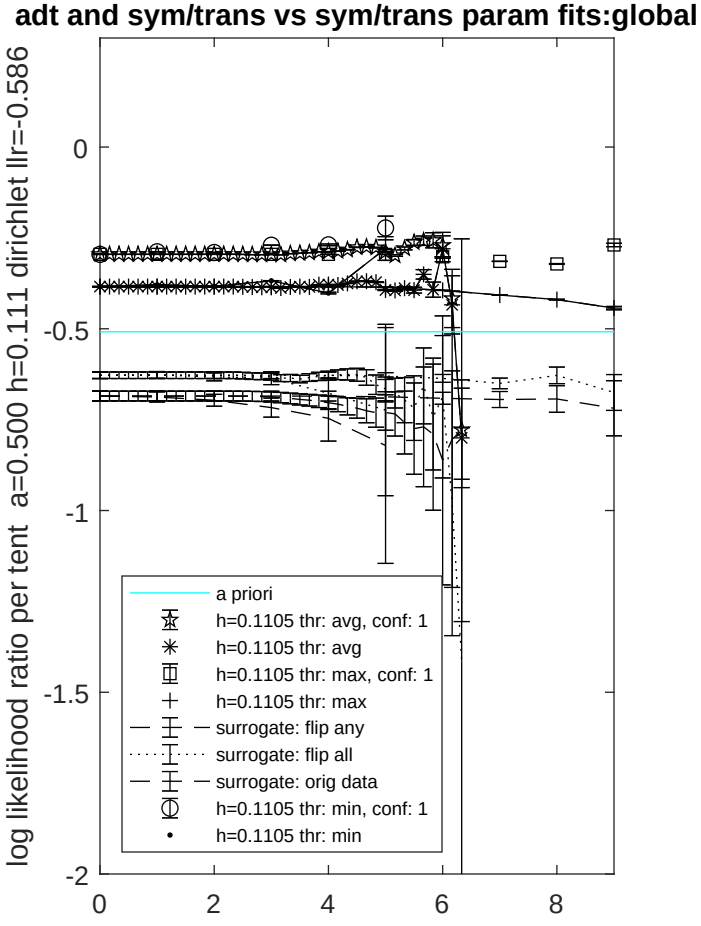
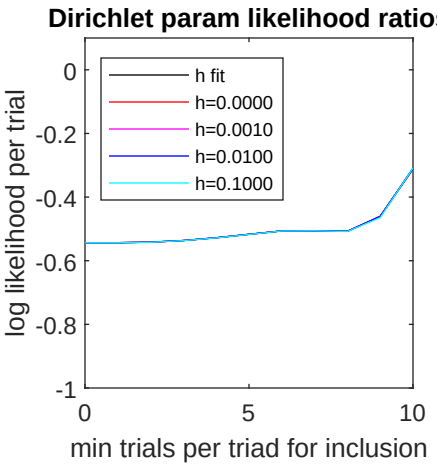
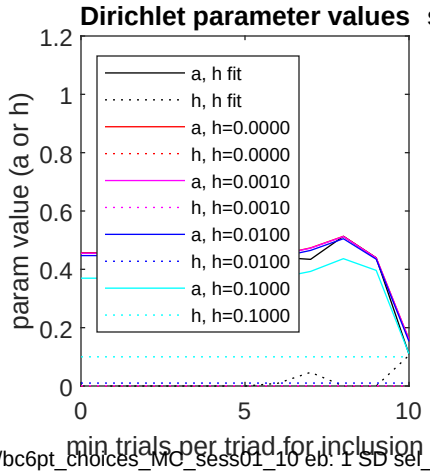
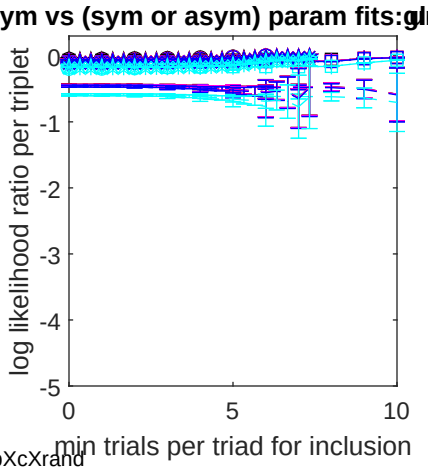
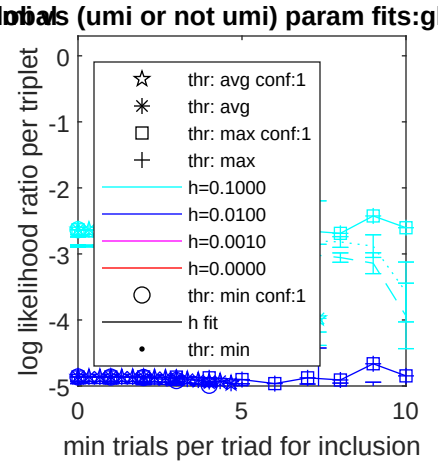

adt and sym/trans vs sym/trans param fits:global
log likelihood ratio per tent a=0.500 h=0.111 dirichlet llr=-0.586
0
-0.5
-1
-1.5
-2
0
2
4
6
8
min trials per triad for inclusion
a priori
h=0.1105 thr: avg, conf: 1
h=0.1105 thr: avg
h=0.1105 thr: max, conf: 1
h=0.1105 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.1105 thr: min, conf: 1
h=0.1105 thr: min
./psg_data/bc6pt_choices_BL_sess01_10 eb: 1 SD, fitted n sel_bmXcmXrand_a05









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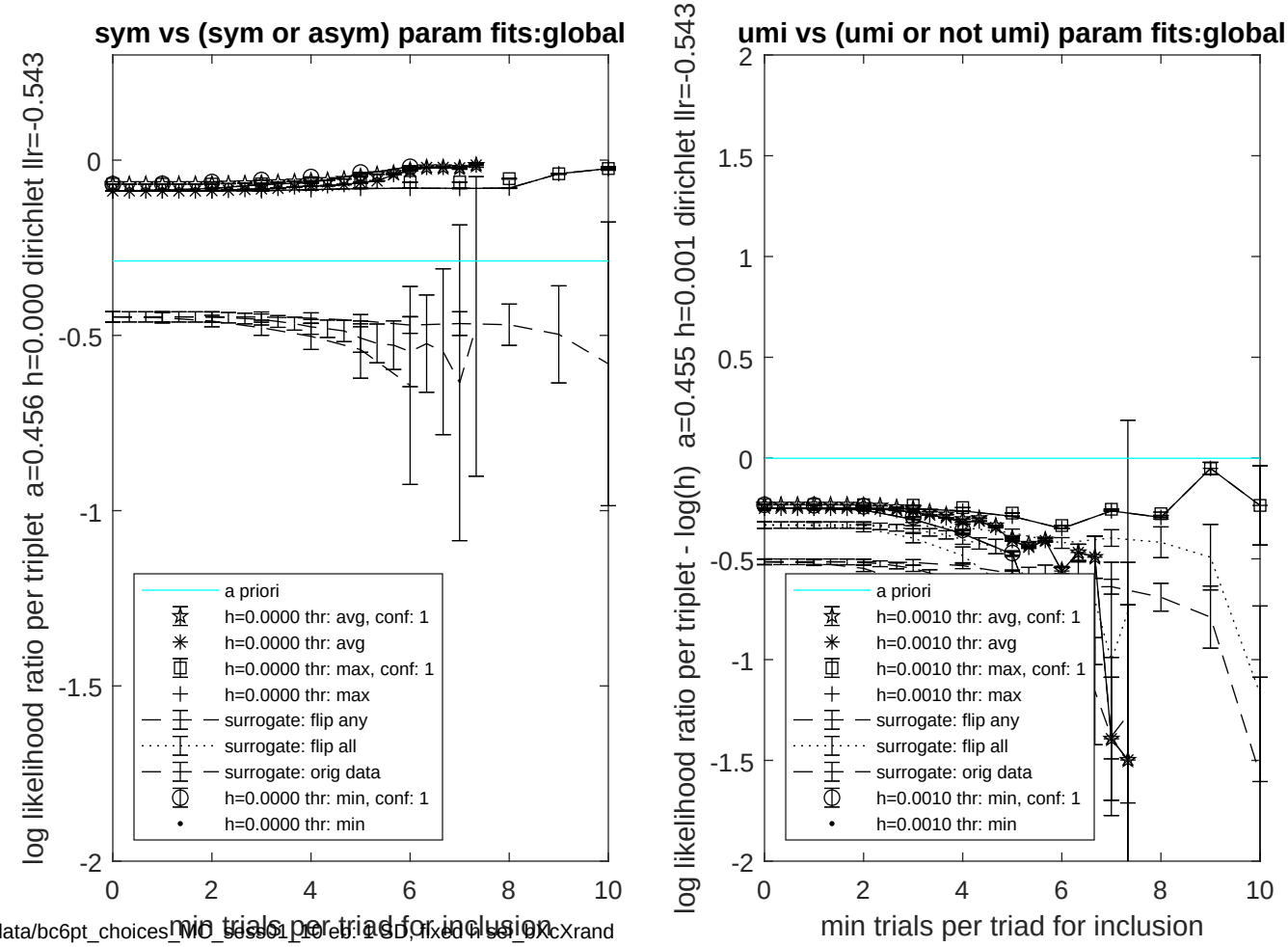

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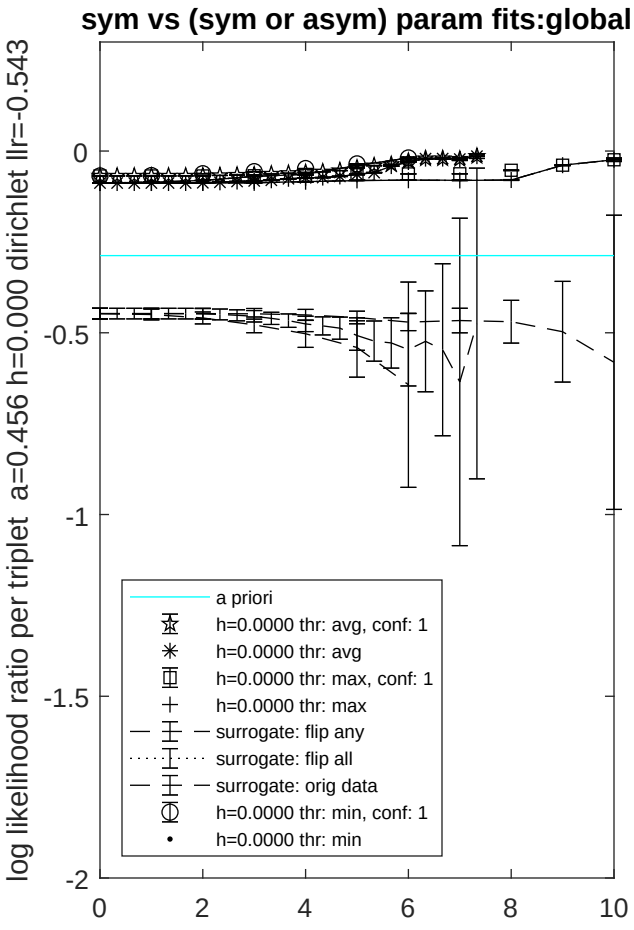

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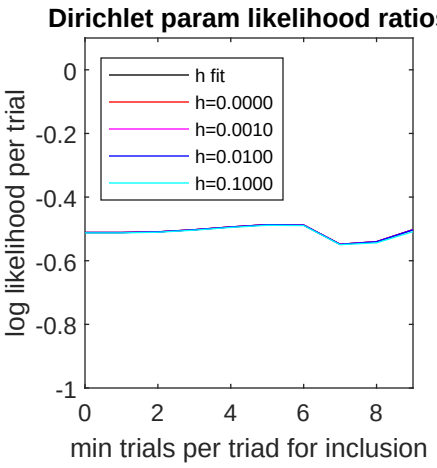

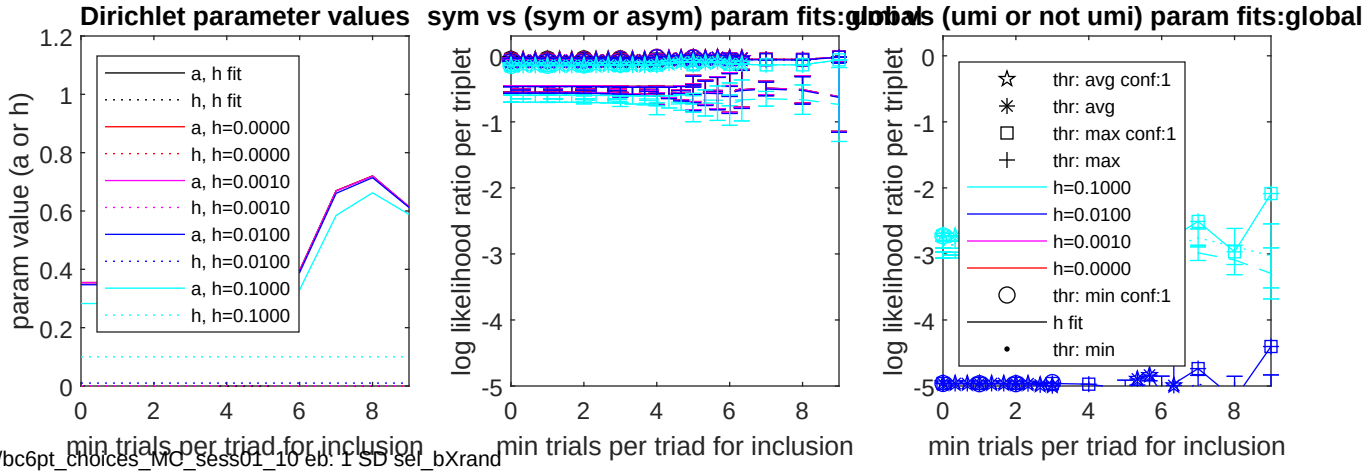

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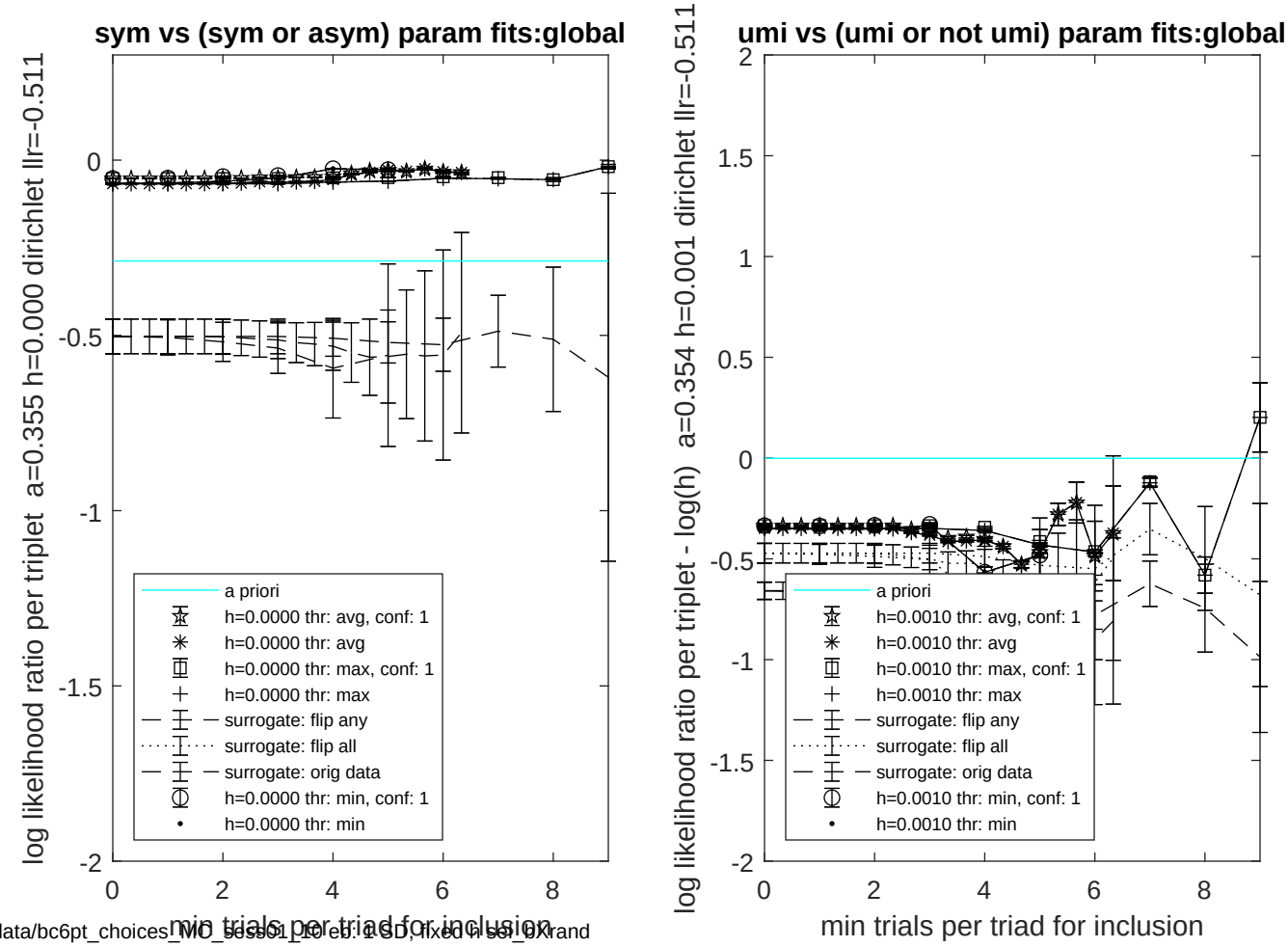

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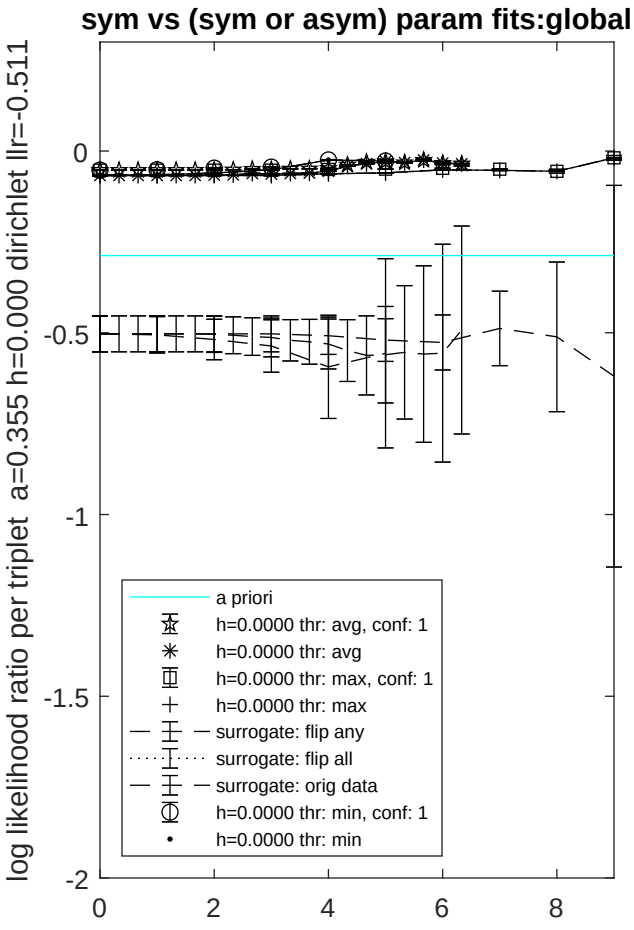

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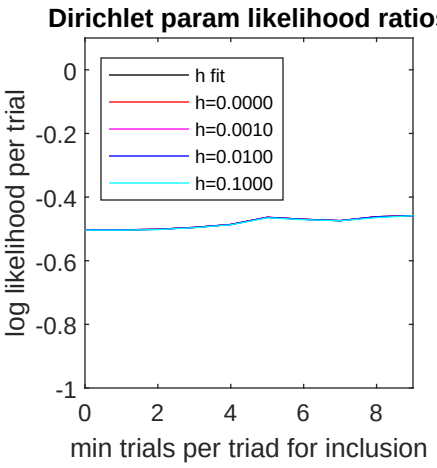

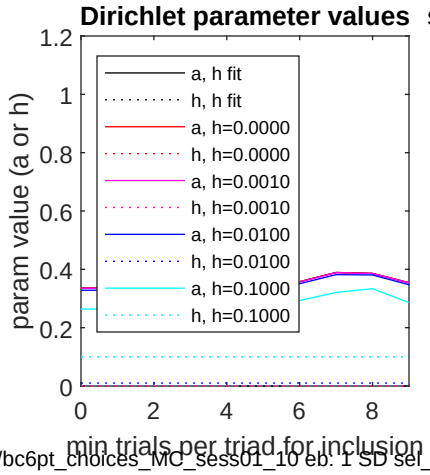

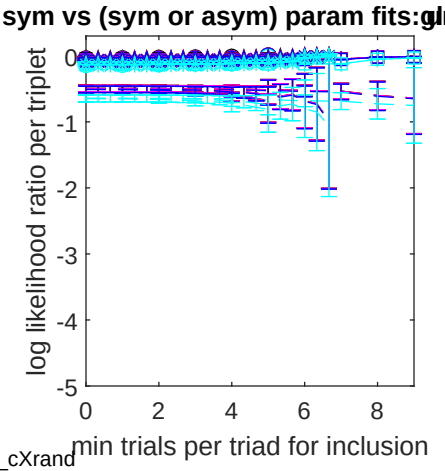

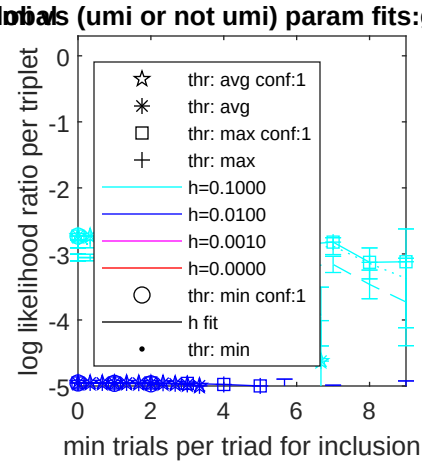

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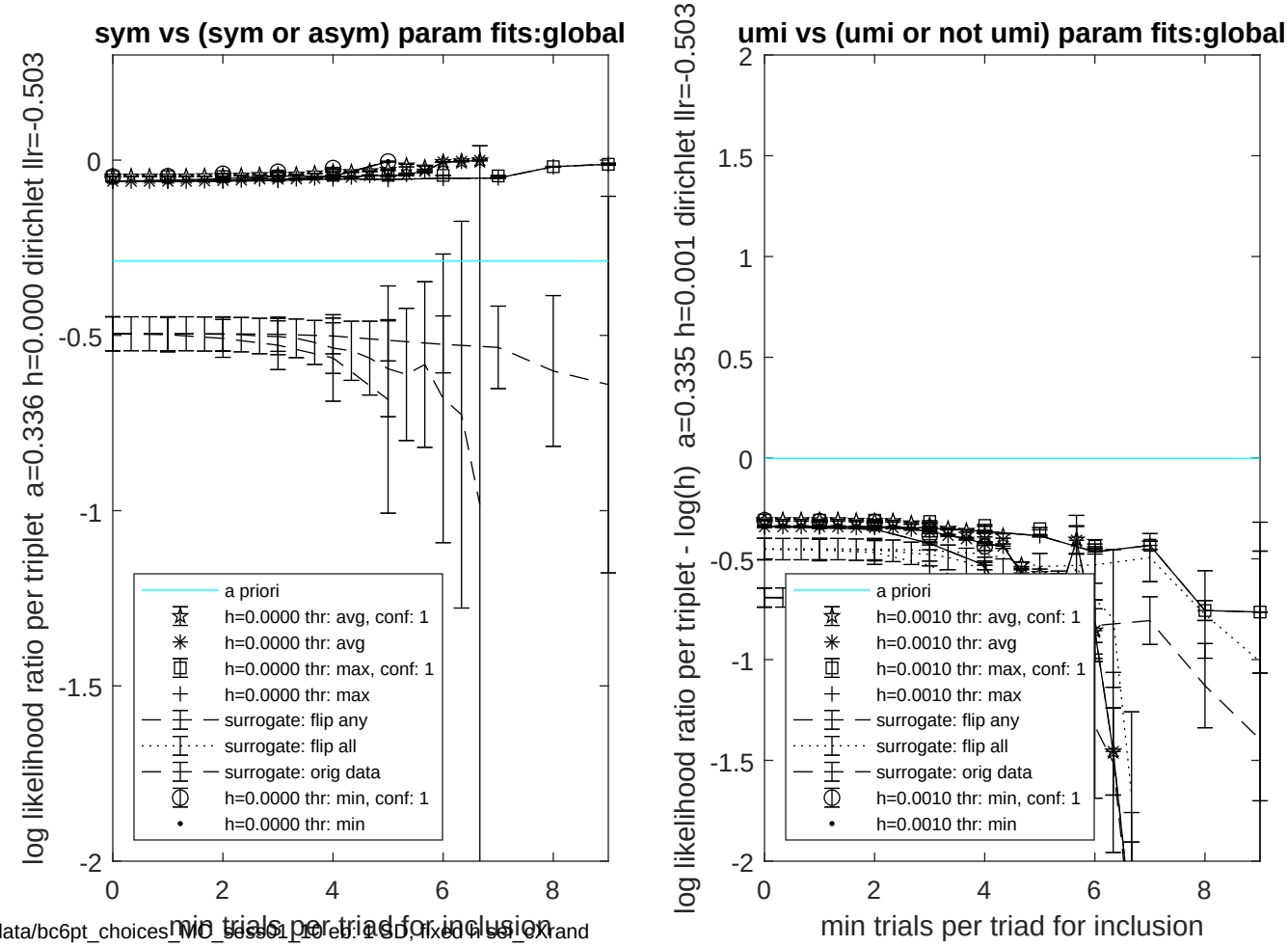

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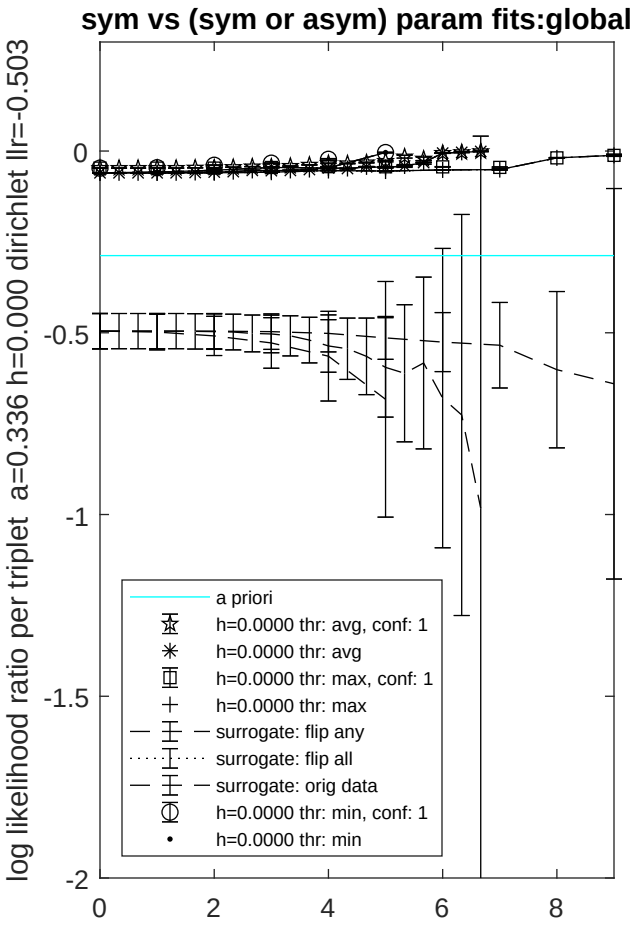

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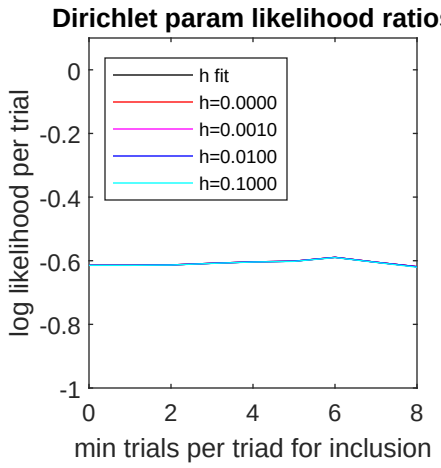
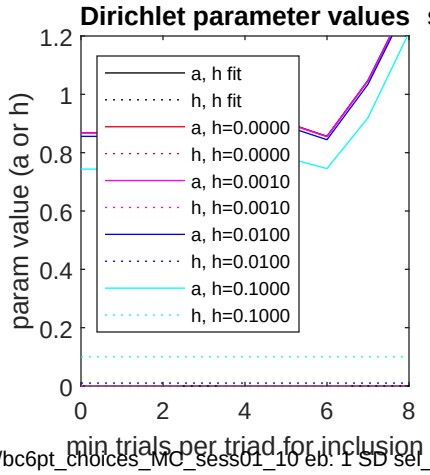
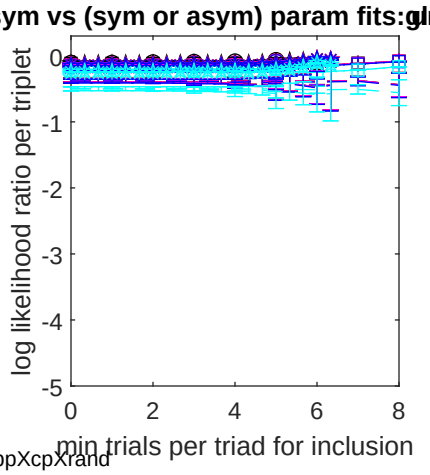
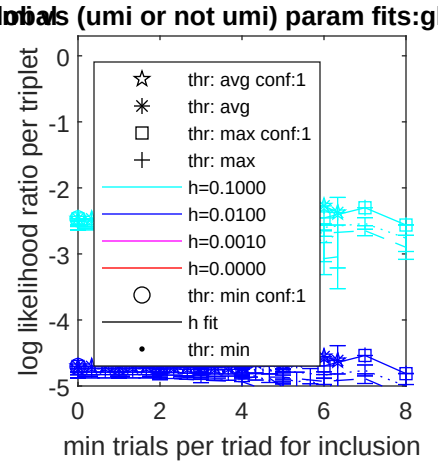
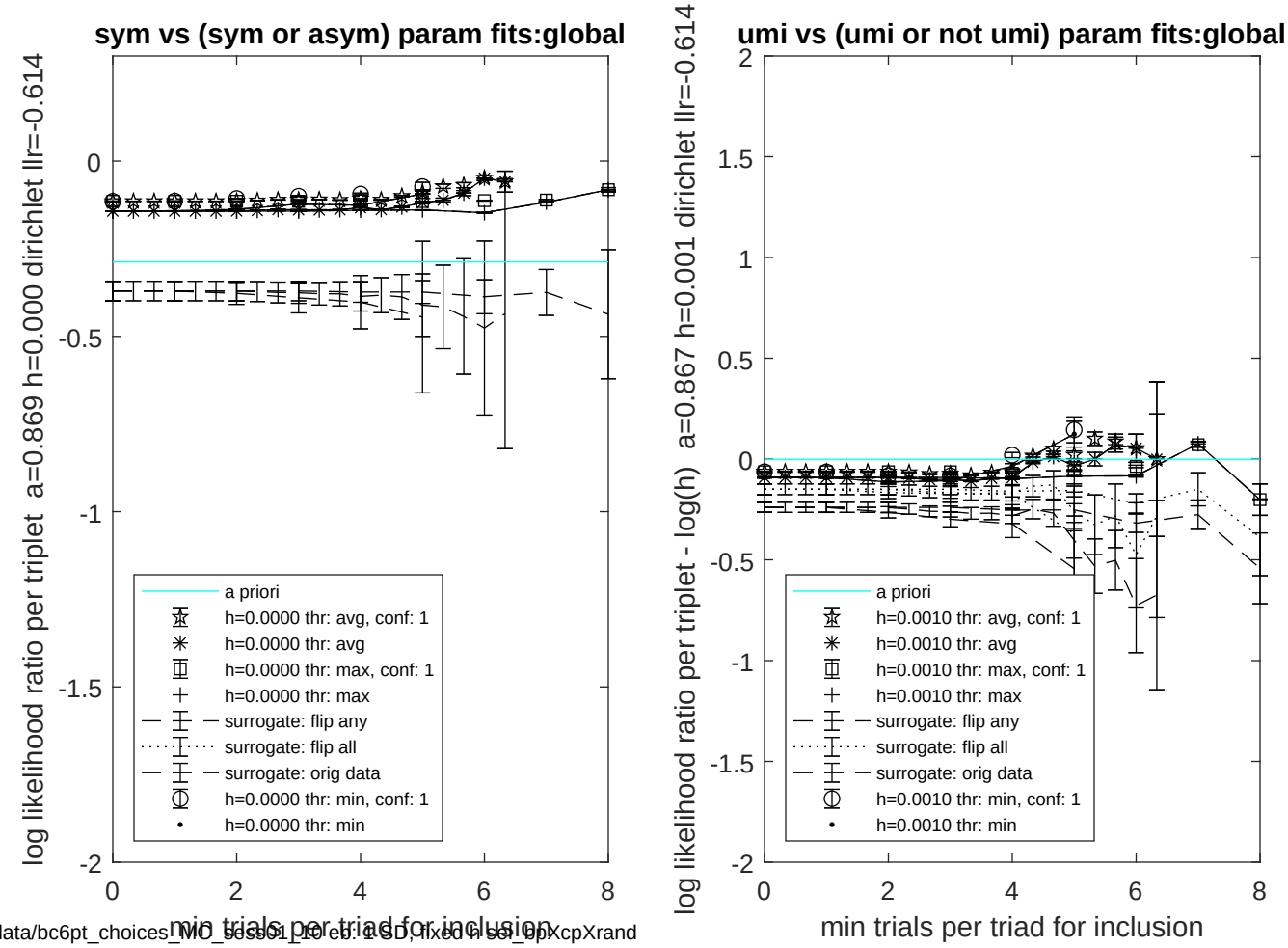




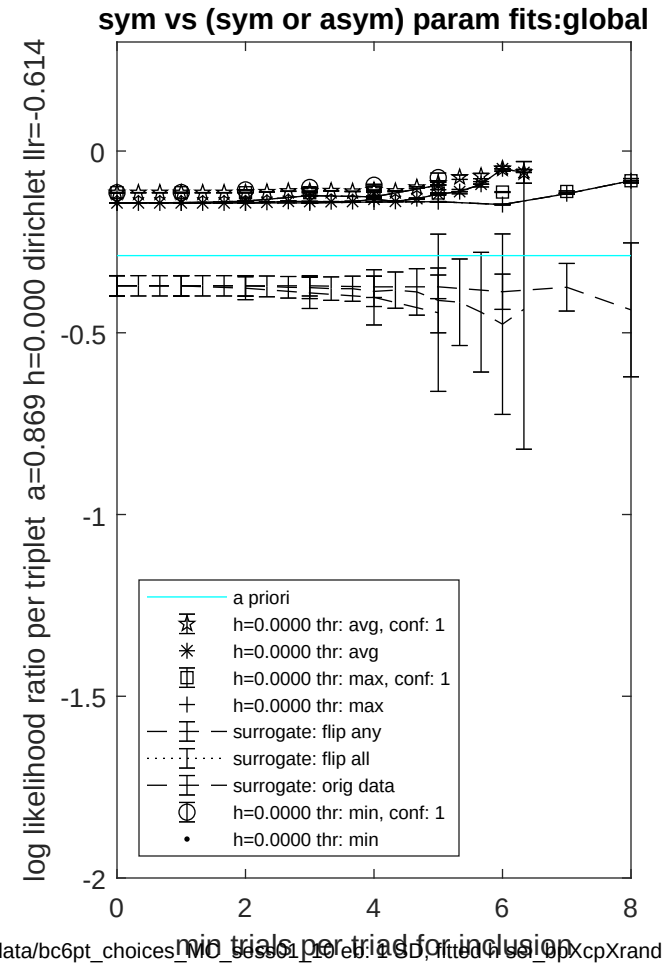


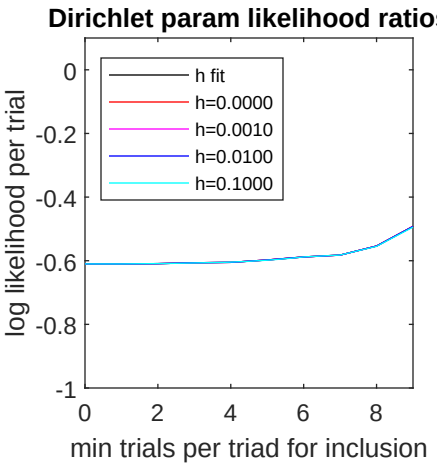


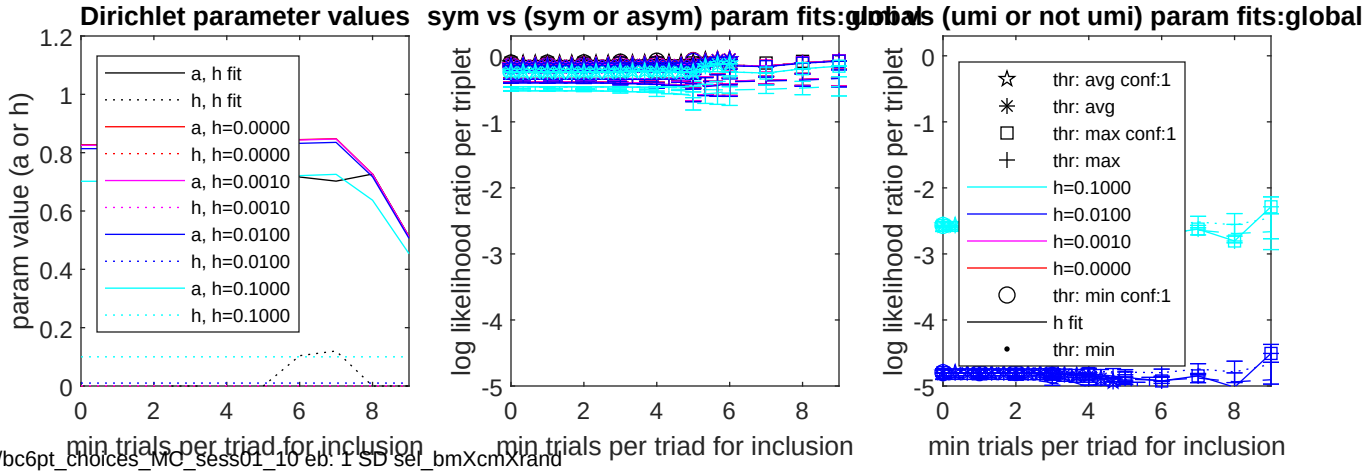
./psg_data/bc6pt_choices_MC_sess01_10 eb: 1 SD sel_bpXcpXrand



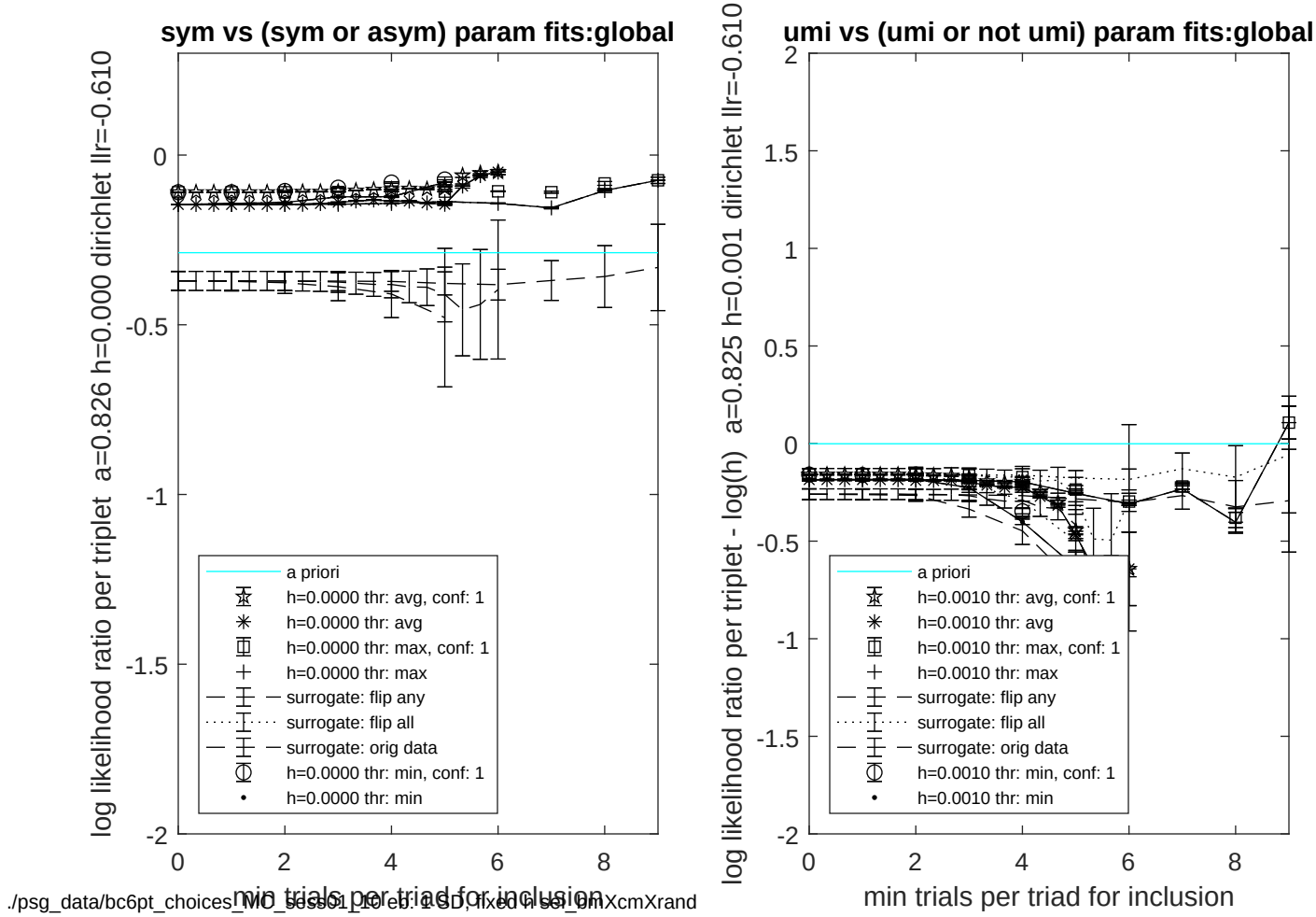
./psg_data/bc6pt_choices_MO_sess01_10 eb: 1 SD, fixed h set_bpXcpXrand

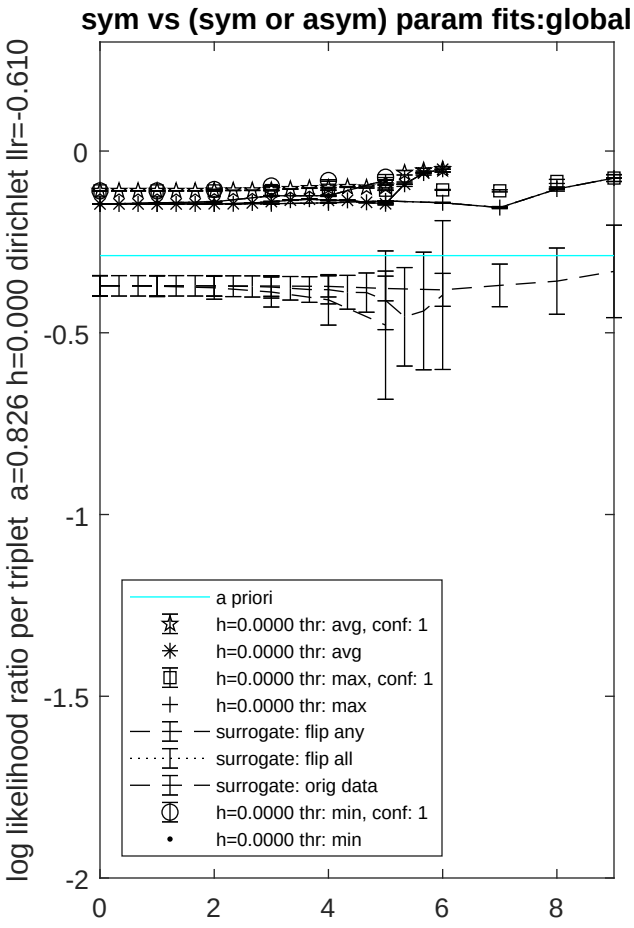

sym vs (sym or asym) param fits:global
log likelihood ratio per triplet a=0.869 h=0.000 dirichlet llr=-0.614
0
-0.5
-1
-1.5
-2
0
2
4
6
8
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
./psg_data/bc6pt_choices_MC_sess01_10 ed: 1 SD, fitted h sel_bpXcpXrand

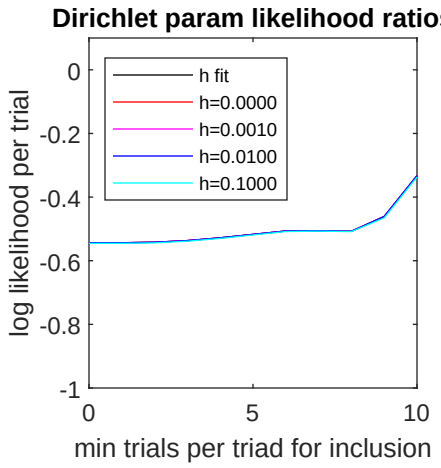


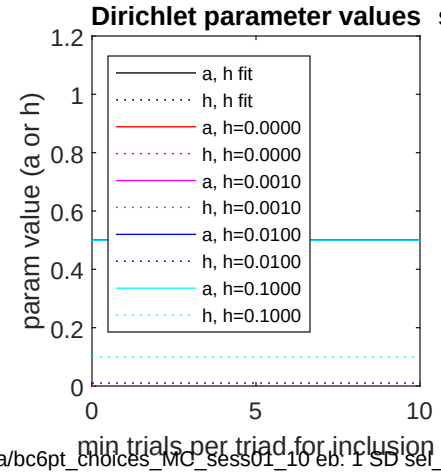


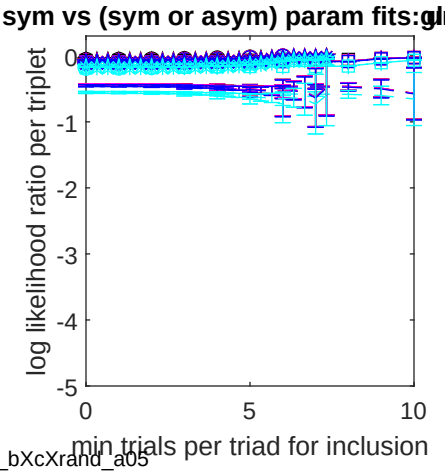
./psg_data/bc6pt_choices_MC_sess01_10 eb: 1 SD sel_bmXcmXrand

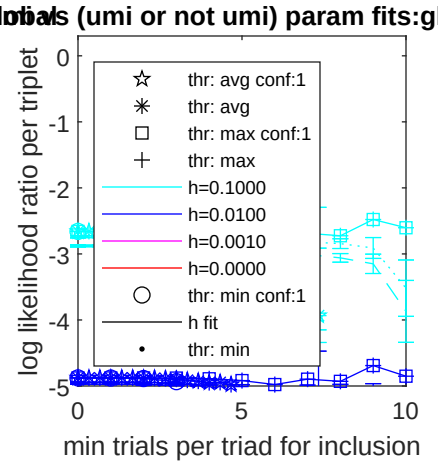

sym vs (sym or asym) param fits:global
log likelihood ratio per triplet a=0.826 h=0.000 dirichlet llr=-0.610
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
./psg_data/bc6pt_choices_MO_sess01_10 eb: 1 SD, fixed h set_bmXcmXrand
umi vs (umi or not umi) param fits:global
log likelihood ratio per triplet - log(h) a=0.825 h=0.001 dirichlet llr=-0.610
min trials per triad for inclusion
a priori
h=0.0010 thr: avg, conf: 1
h=0.0010 thr: avg
h=0.0010 thr: max, conf: 1
h=0.0010 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0010 thr: min, conf: 1
h=0.0010 thr: min


sym vs (sym or asym) param fits:global
log likelihood ratio per triplet a=0.826 h=0.000 dirichlet llr=-0.610
0
-0.5
-1
-1.5
-2
0
2
4
6
8
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
./psg_data/bc6pt_choices_MO_sess01_10 eb: 1 SD, fitted h sel_bmXcmXrand

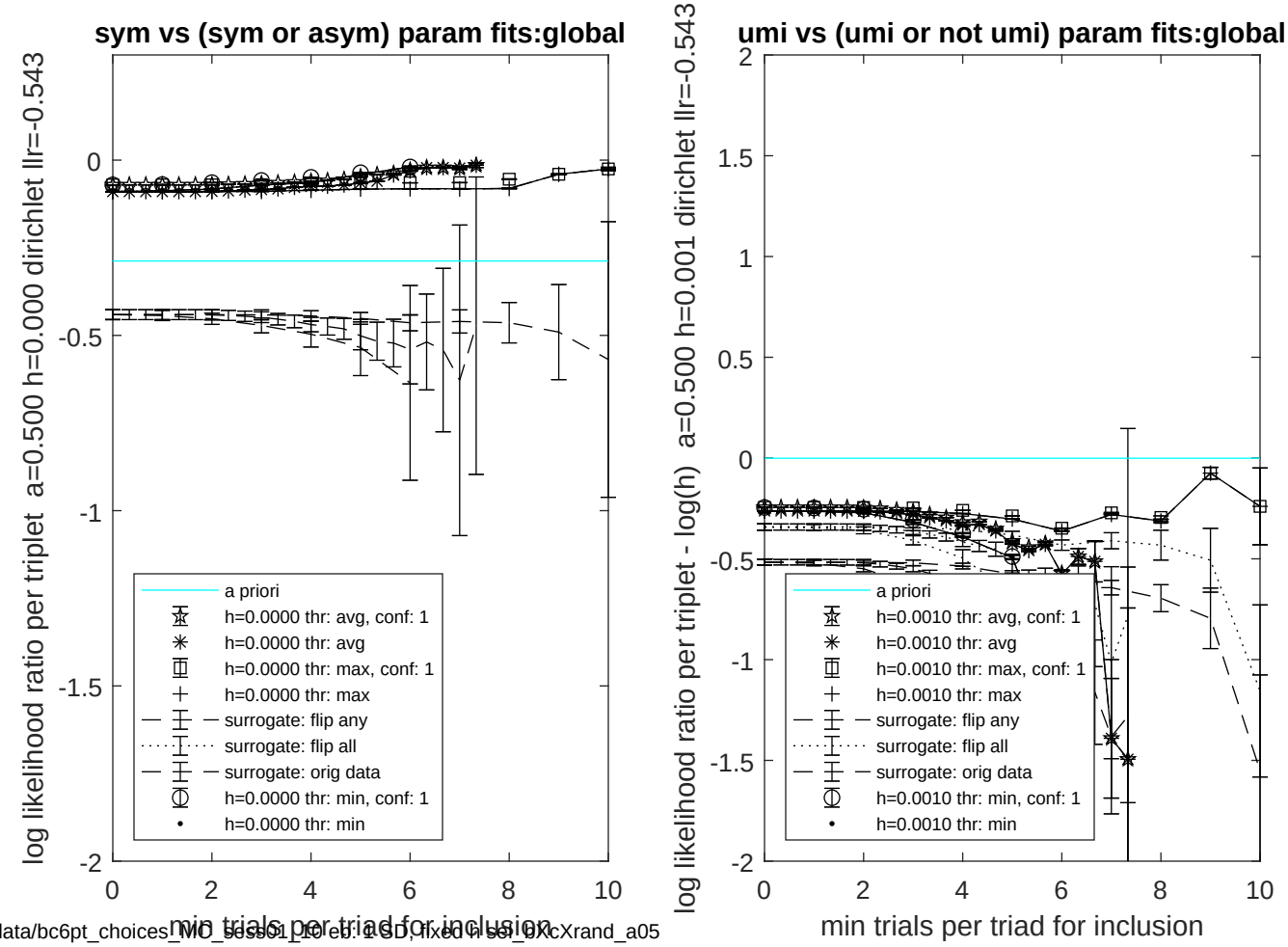




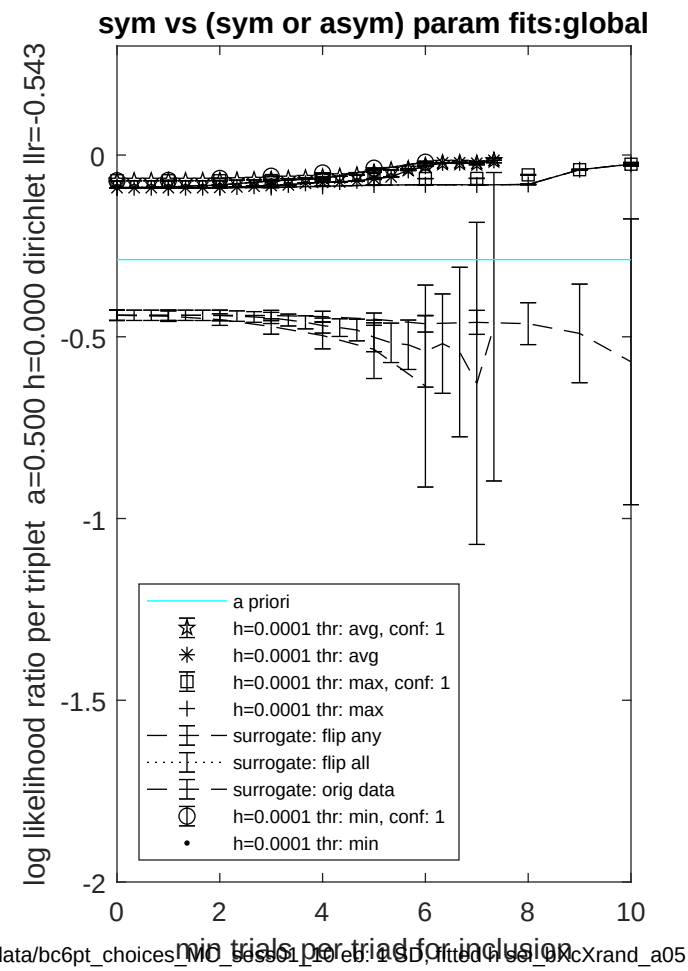


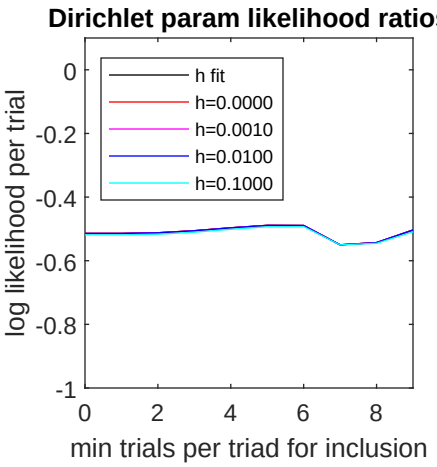


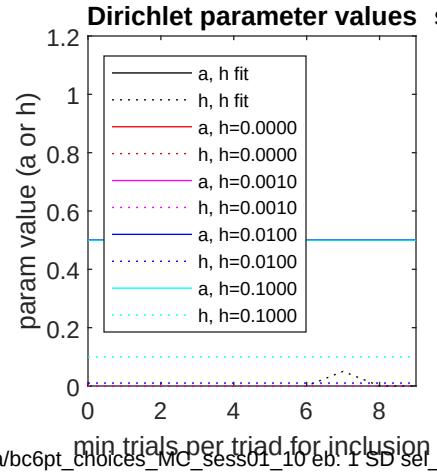
./psg_data/bc6pt_choices_MC_sess01_10 eb: 1 SD sel_bXcXrand_a05

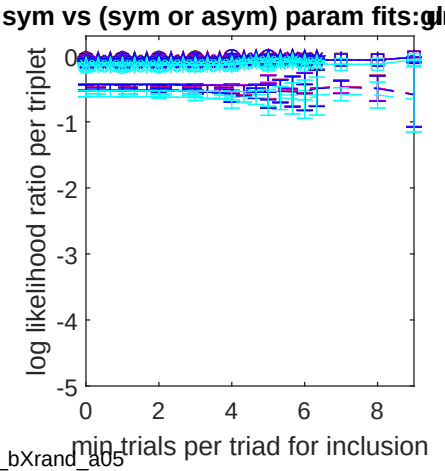


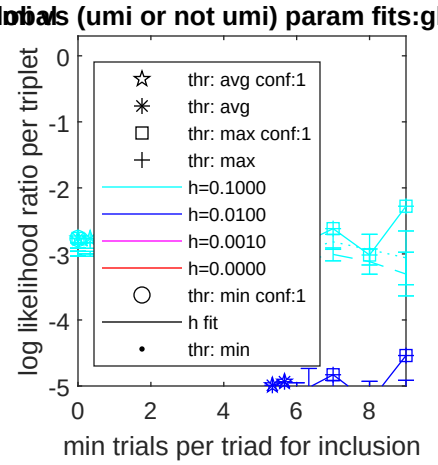
./psg_data/bc6pt_choices_MO_sess01_10 eb: 1 SD, fixed h set_bXcXrand_a05


sym vs (sym or asym) param fits:global
log likelihood ratio per triplet a=0.500 h=0.000 dirichlet llr=-0.543
0
-0.5
-1
-1.5
-2
0
2
4
6
8
10
min trials per triad for inclusion
a priori
h=0.0001 thr: avg, conf: 1
h=0.0001 thr: avg
h=0.0001 thr: max, conf: 1
h=0.0001 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0001 thr: min, conf: 1
h=0.0001 thr: min
./psg_data/bc6pt_choices_MC_sess01_10 eb: 1 SD, fitted h sel_bXcXrand_a05









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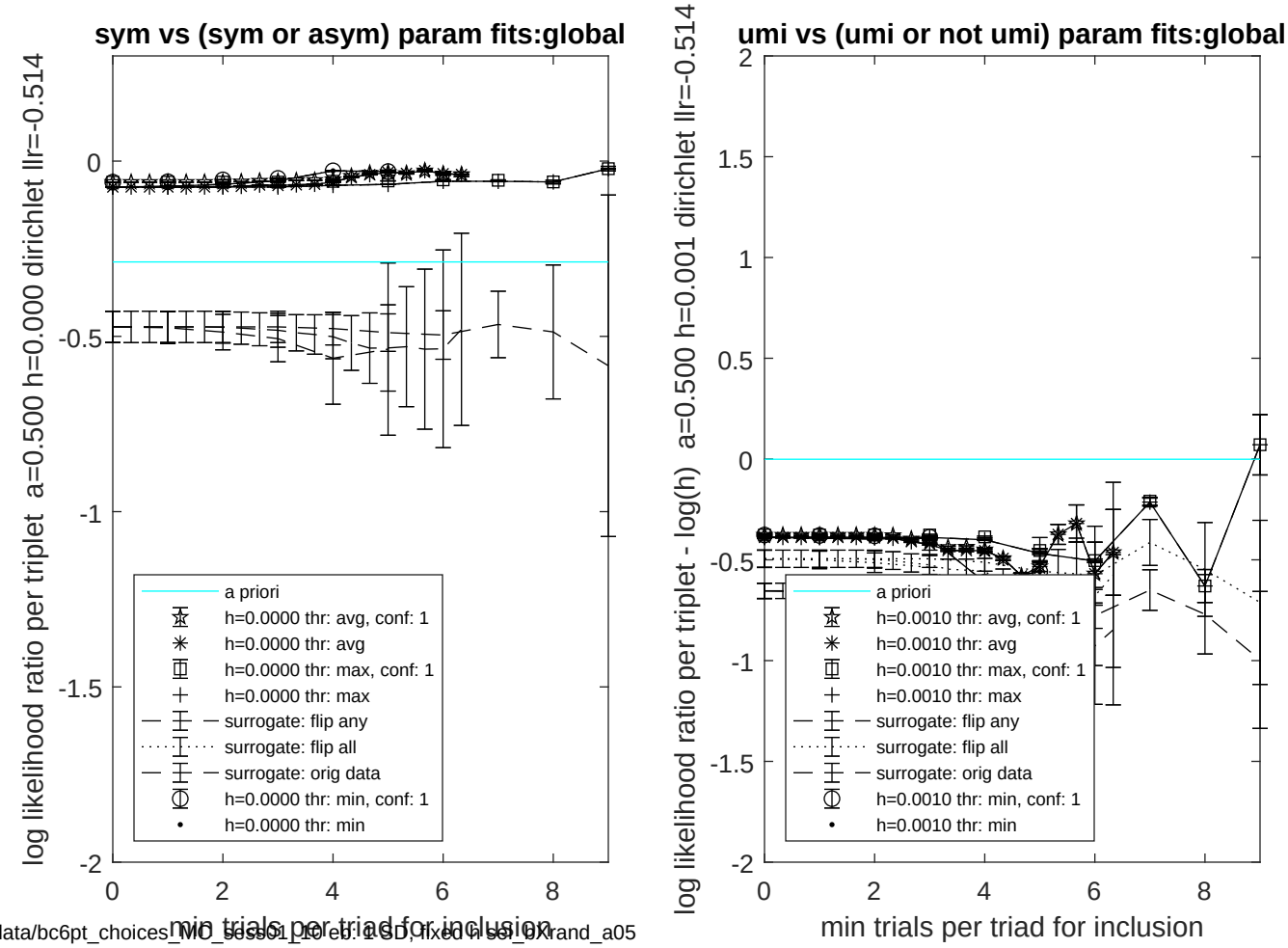

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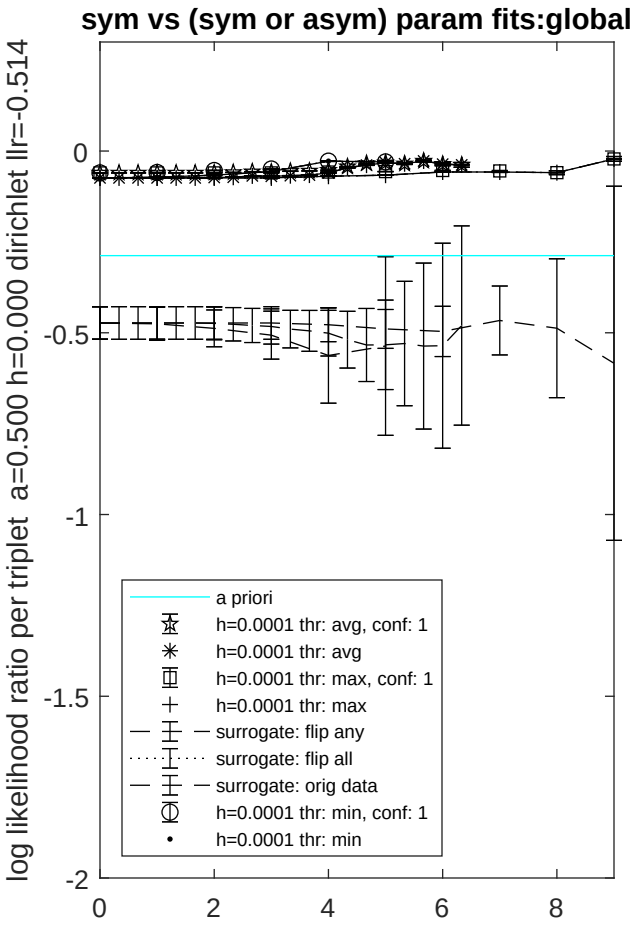
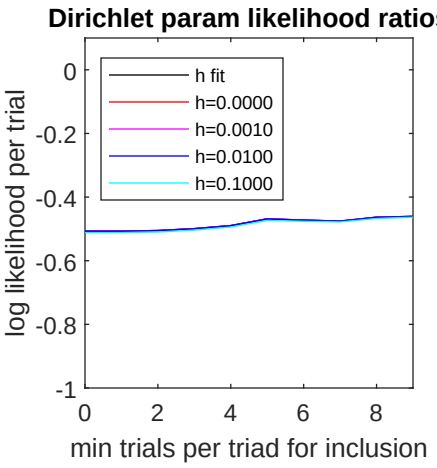
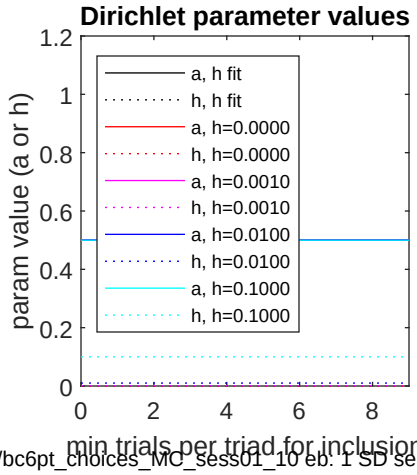
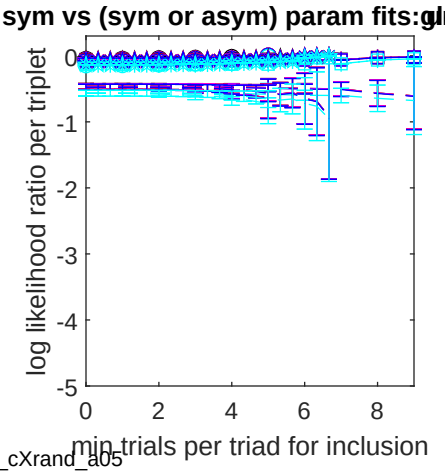
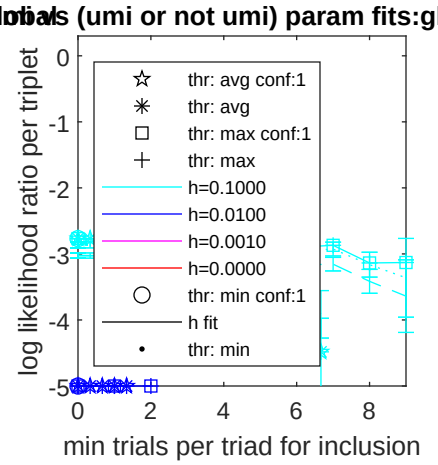

sym vs (sym or asym) param fits:global
log likelihood ratio per triplet a=0.500 h=0.000 dirichlet llr=-0.514
0
-0.5
-1
-1.5
-2
0
2
4
6
8
min trials per triad for inclusion
a priori
h=0.0001 thr: avg, conf: 1
h=0.0001 thr: avg
h=0.0001 thr: max, conf: 1
h=0.0001 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0001 thr: min, conf: 1
h=0.0001 thr: min
./psg_data/bc6pt_choices_MC_sess01_10 eb: 1 SD, fitted h sel_bXrand_a05









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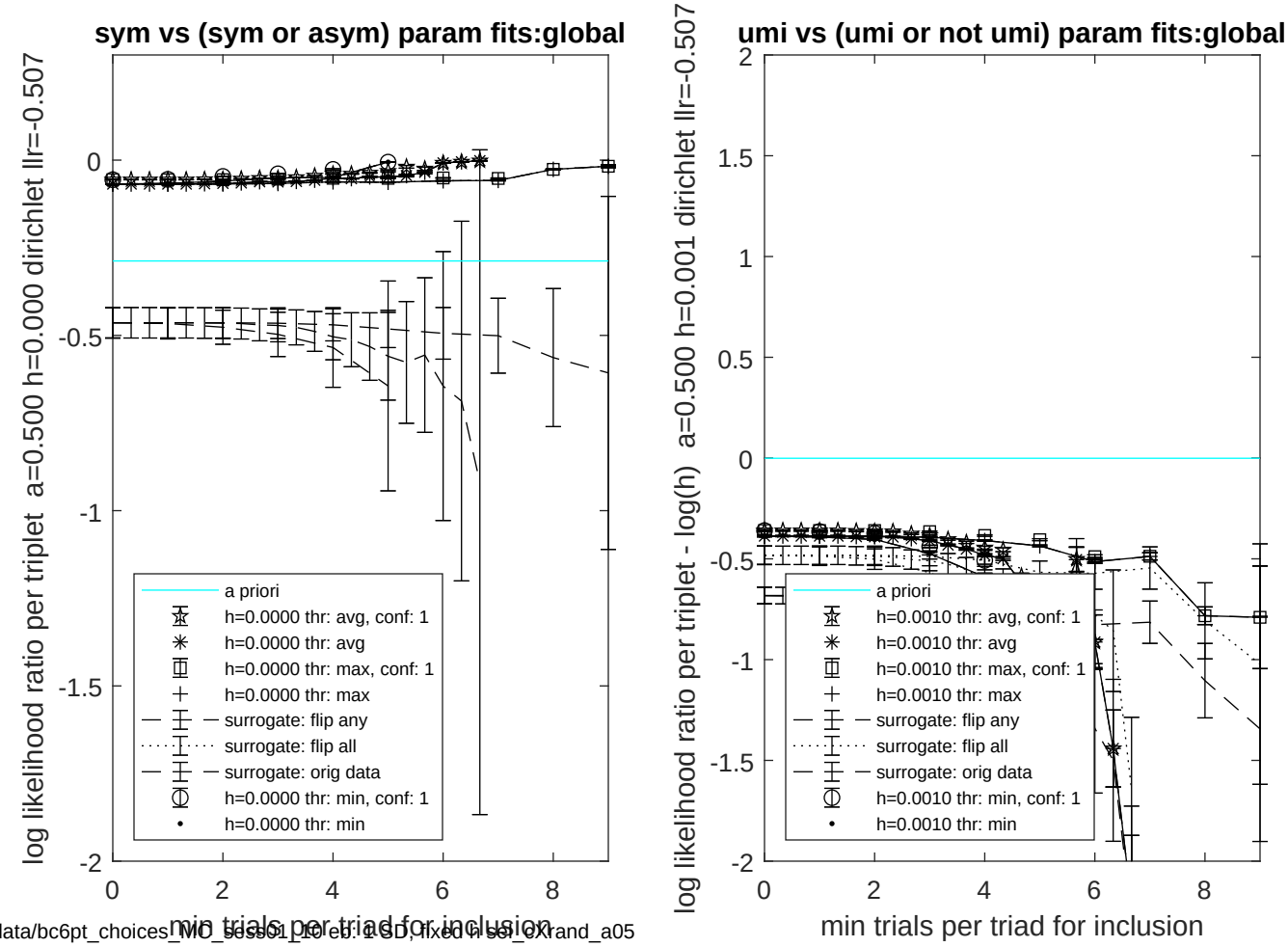

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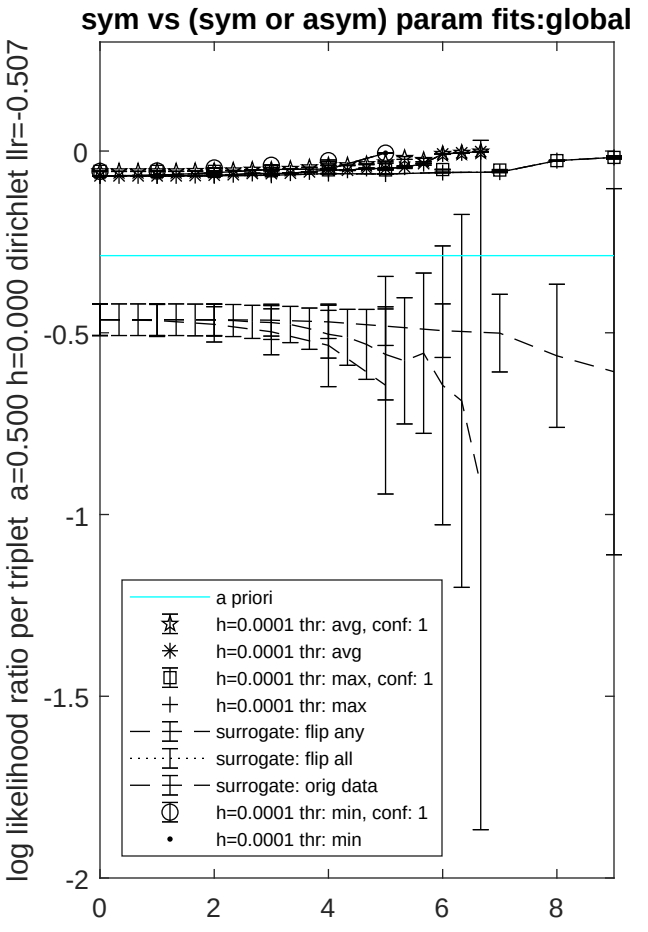
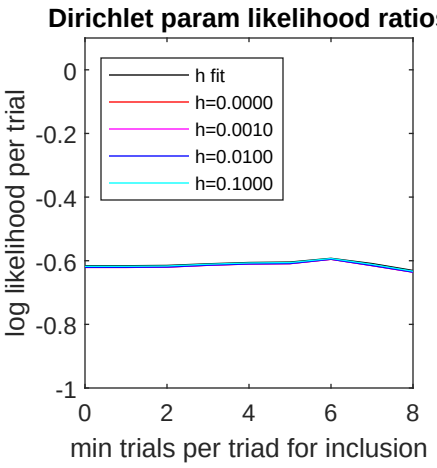
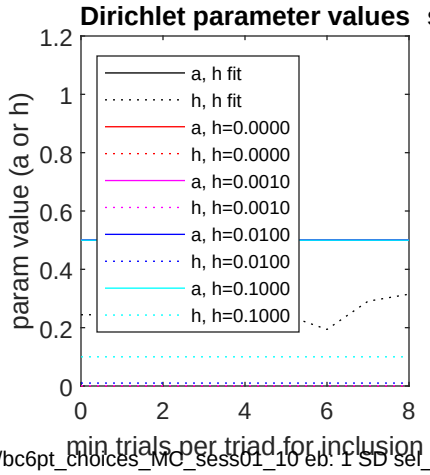
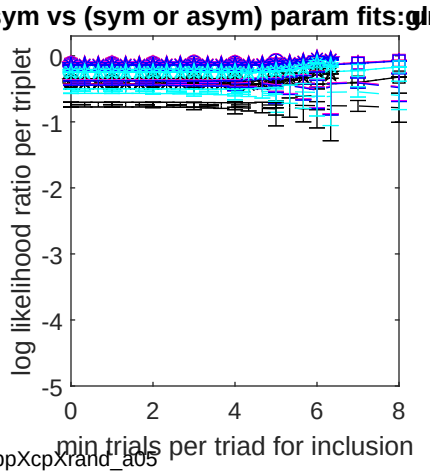
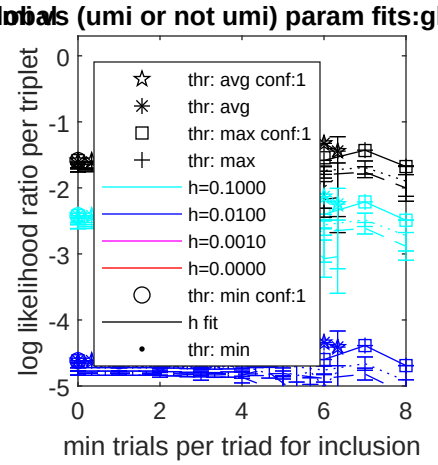

sym vs (sym or asym) param fits:global
log likelihood ratio per triplet a=0.500 h=0.000 dirichlet llr=-0.507
0
-0.5
-1
-1.5
-2
0
2
4
6
8
min trials per triad for inclusion
a priori
h=0.0001 thr: avg, conf: 1
h=0.0001 thr: avg
h=0.0001 thr: max, conf: 1
h=0.0001 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0001 thr: min, conf: 1
h=0.0001 thr: min
./psg_data/bc6pt_choices_MO_sess01_10 eb: 1 SD, fitted h sel_cXrand_a05









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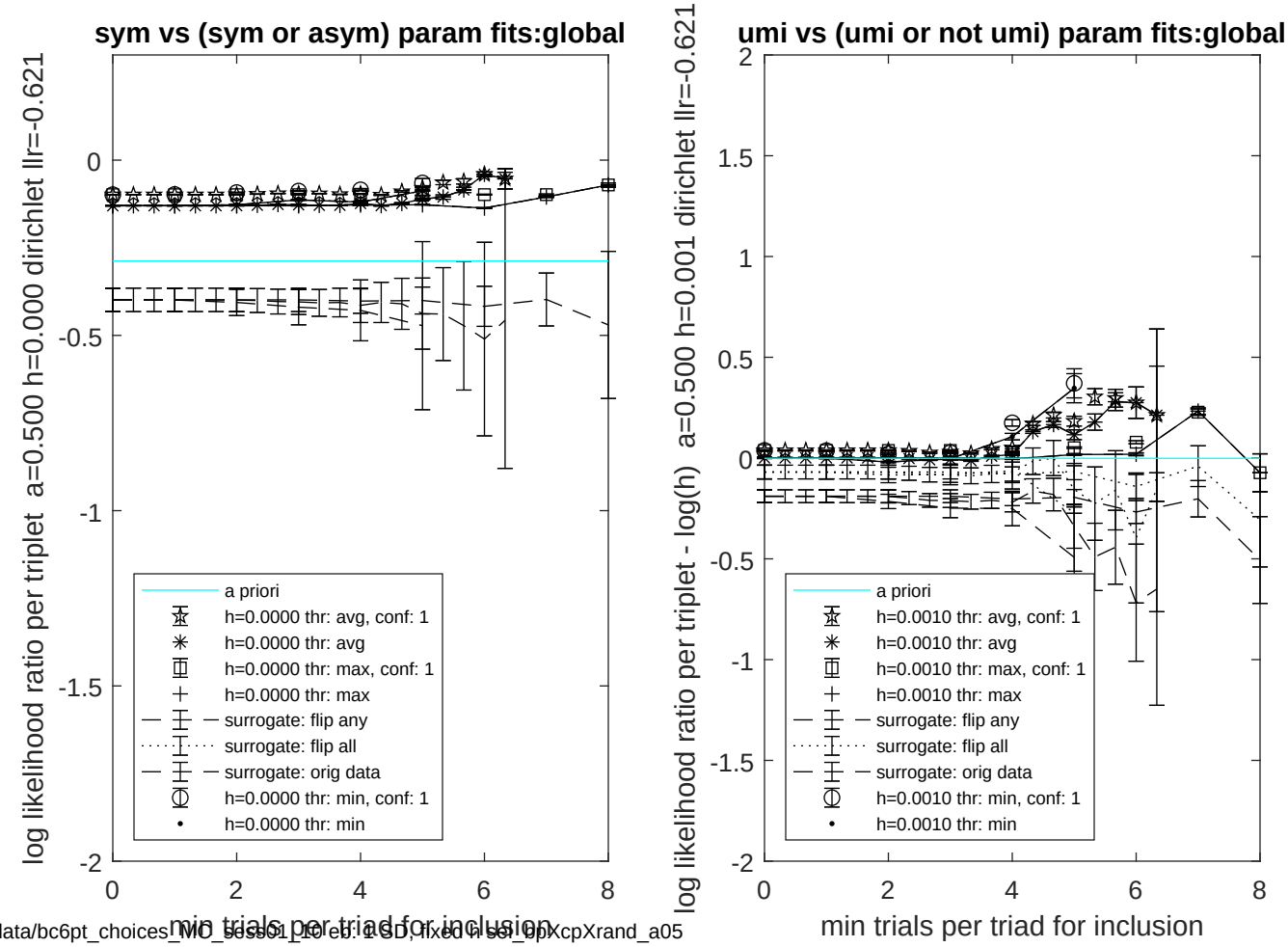

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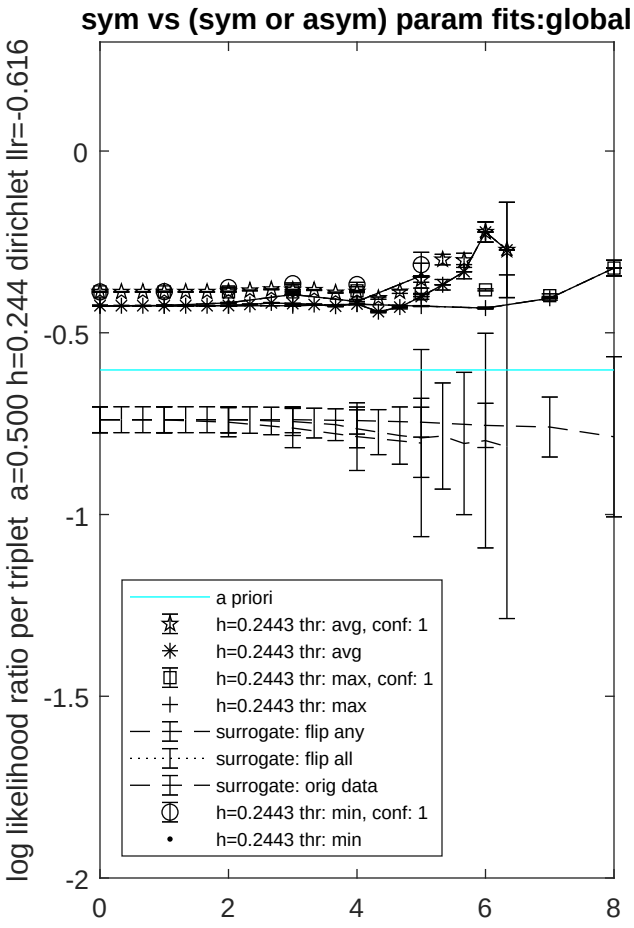

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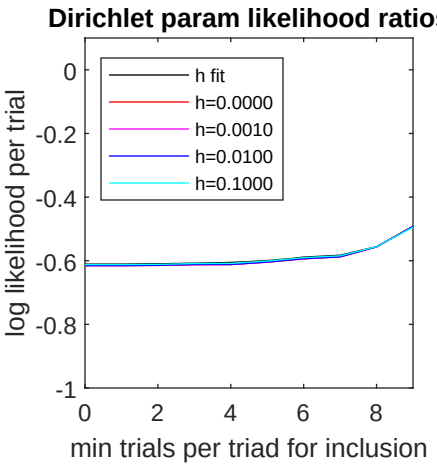

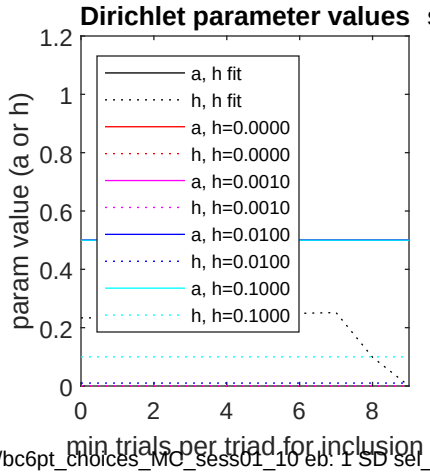

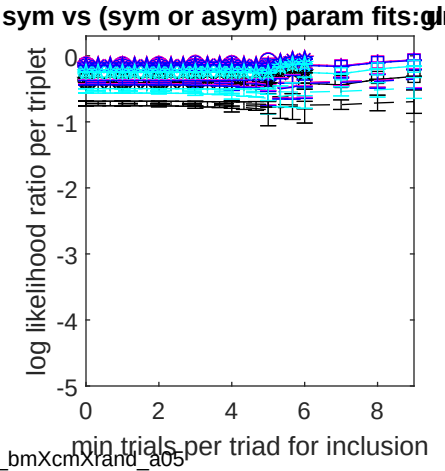

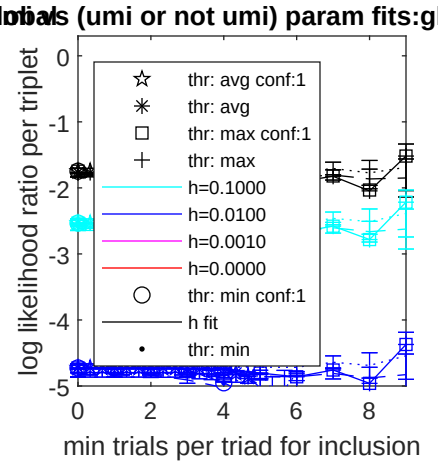

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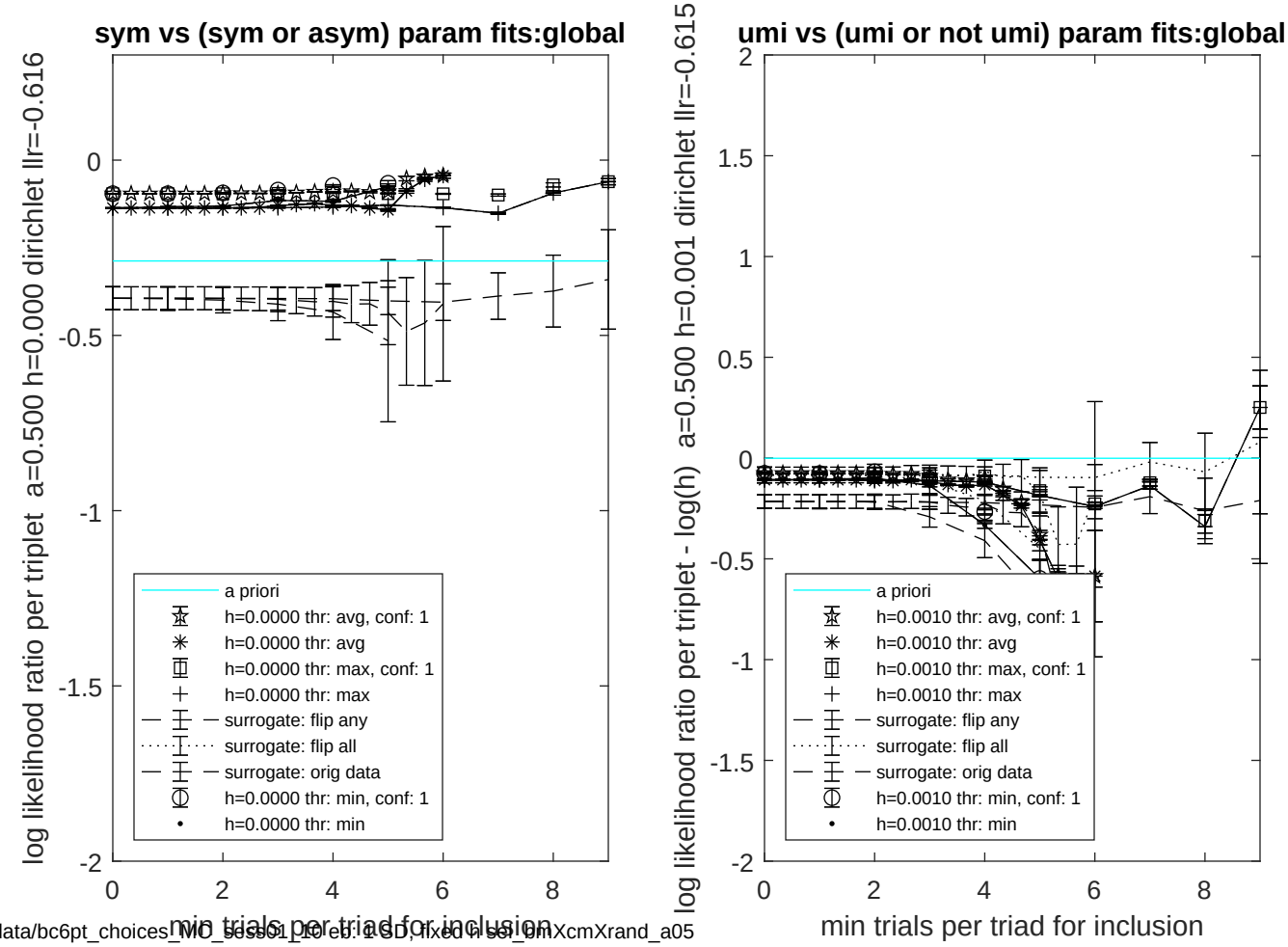

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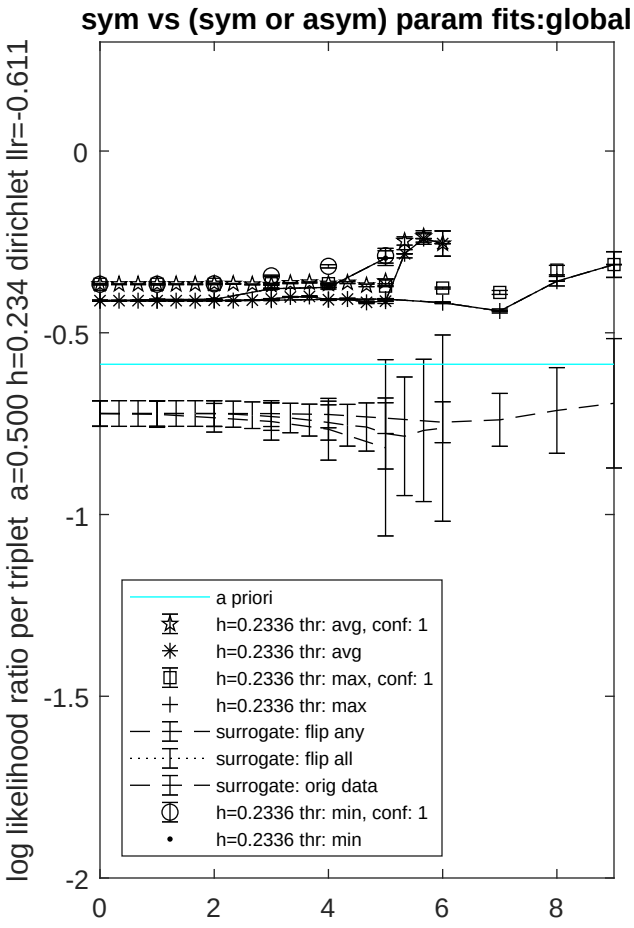

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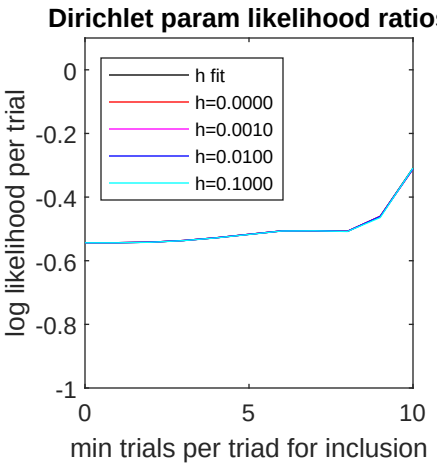

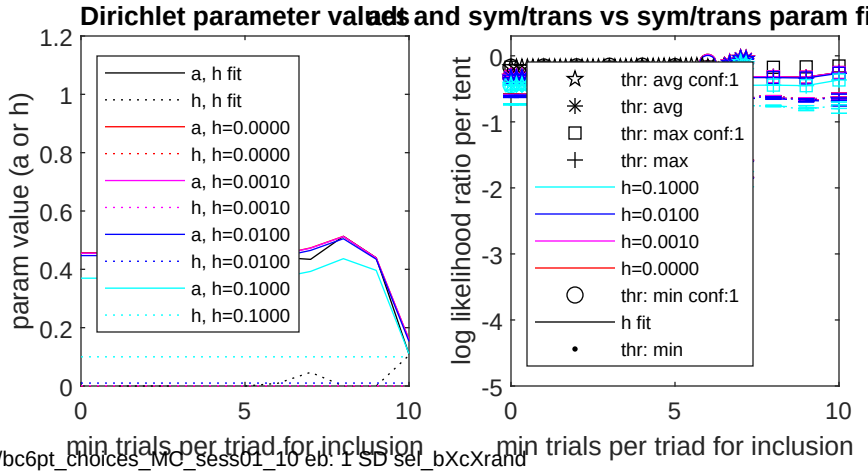

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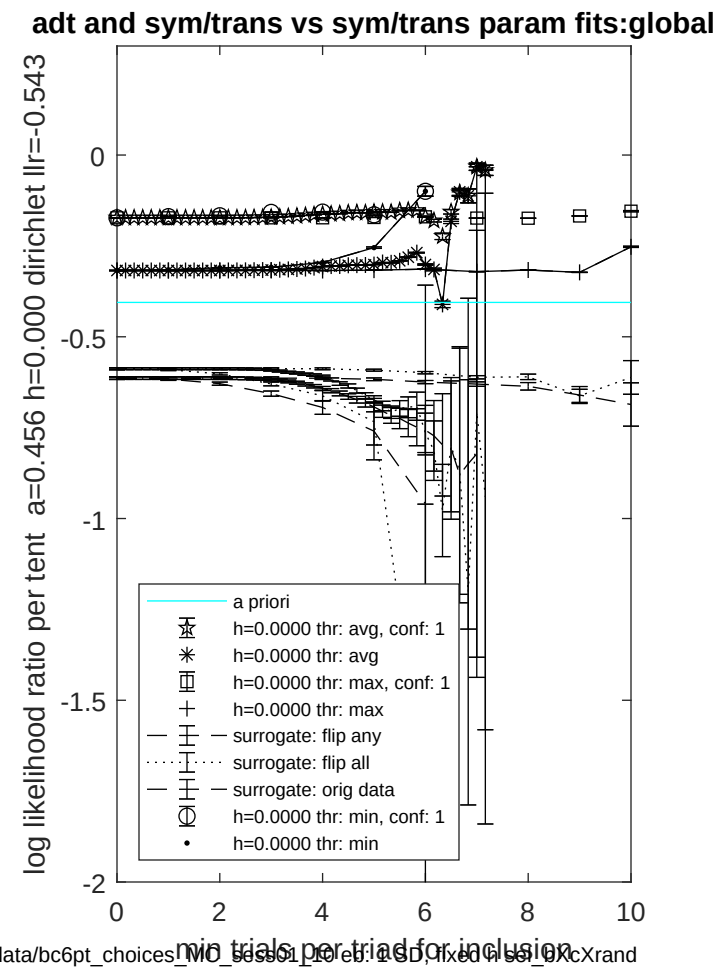
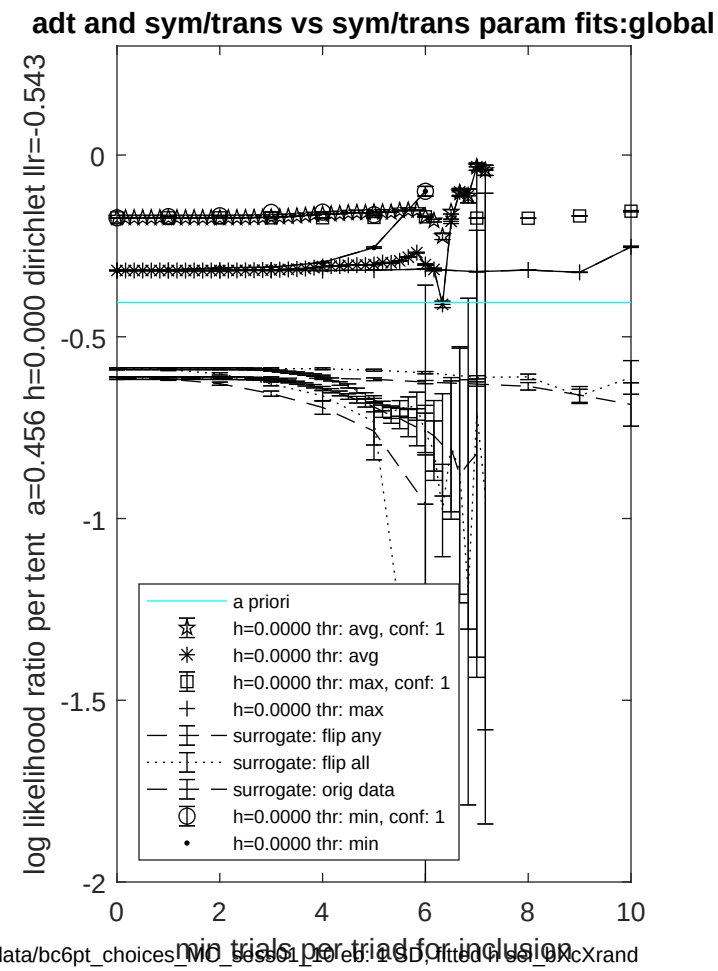
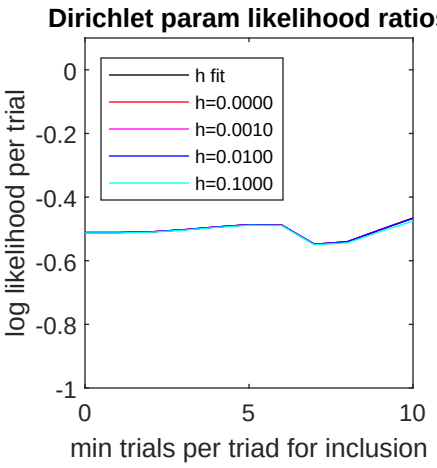
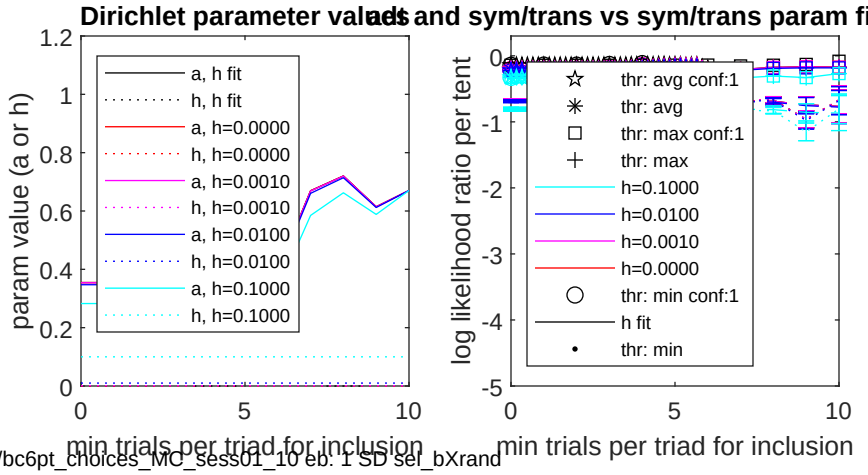
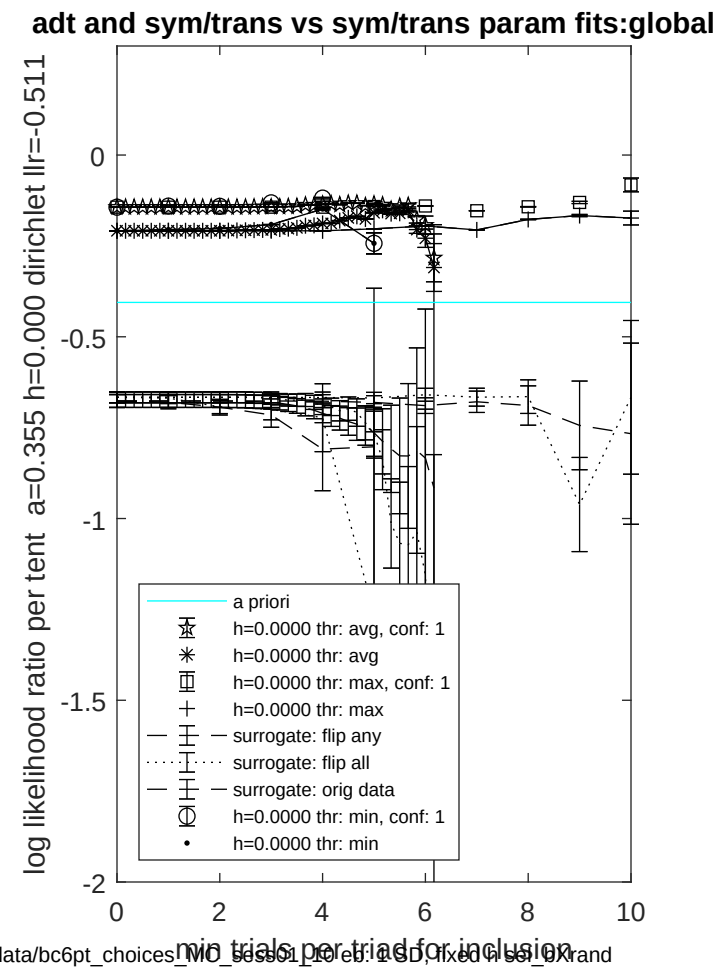

adt and sym/trans vs sym/trans param fits:global
log likelihood ratio per tent a=0.456 h=0.000 dirichlet llr=-0.543
0
-0.5
-1
-1.5
-2
0
2
4
6
8
10
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
./psg_data/bc6pt_choices_MC_sess01_10 eb: 1 SD, fixed h set_bXcXrand


adt and sym/trans vs sym/trans param fits:global
log likelihood ratio per tent a=0.456 h=0.000 dirichlet llr=-0.543
0
-0.5
-1
-1.5
-2
0
2
4
6
8
10
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
./psg_data/bc6pt_choices_MO_sess01_10 eb: 1 SD, fitted h sel_bXcXrand

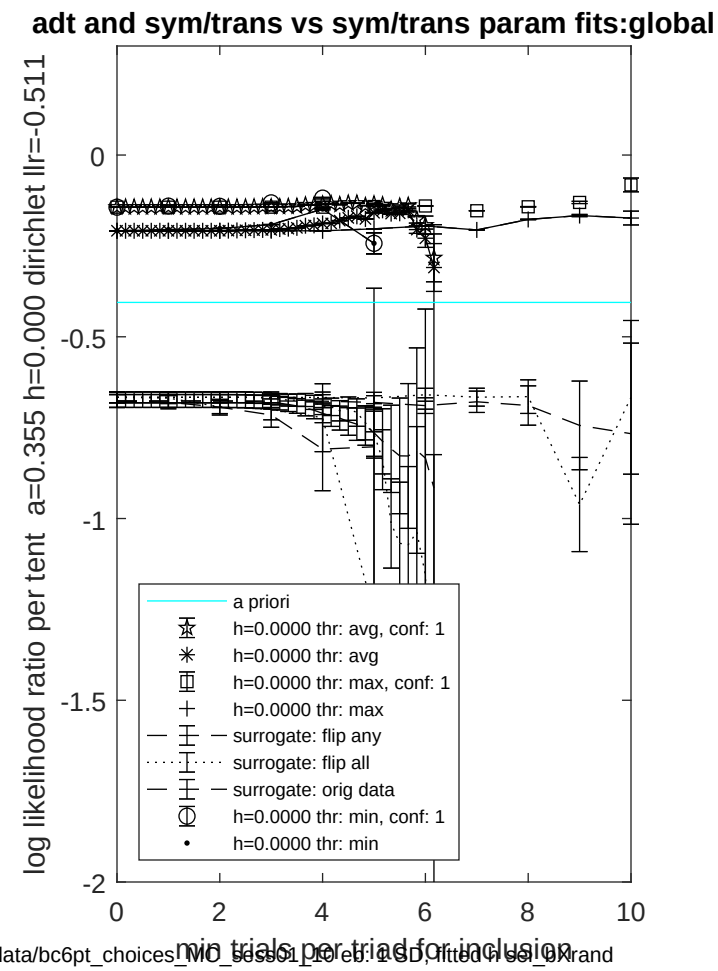




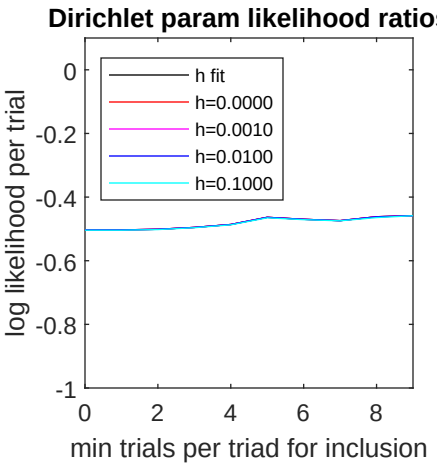
./psg_data/bc6pt_choices_MC_sess01_10 eb: 1 SD sel_bXrand

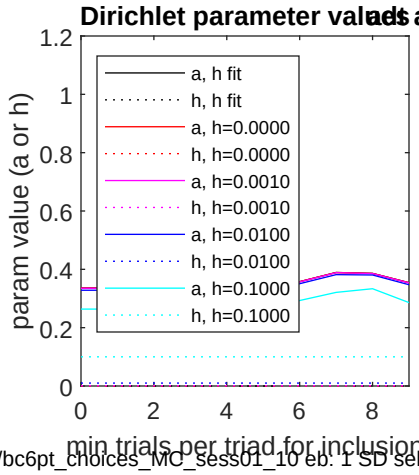


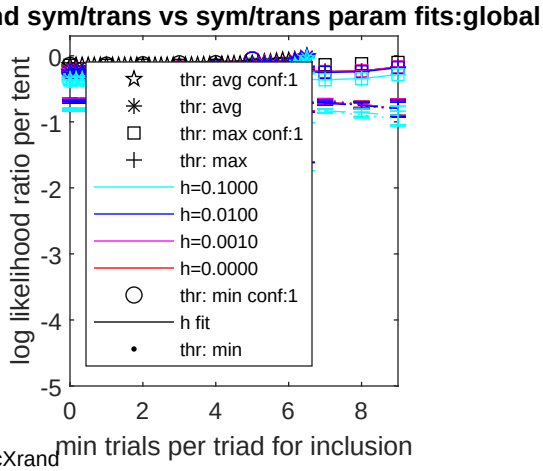
./psg_data/bc6pt_choices_MC_sess01_10 eb: 1 SD, fixed h set_bXrand



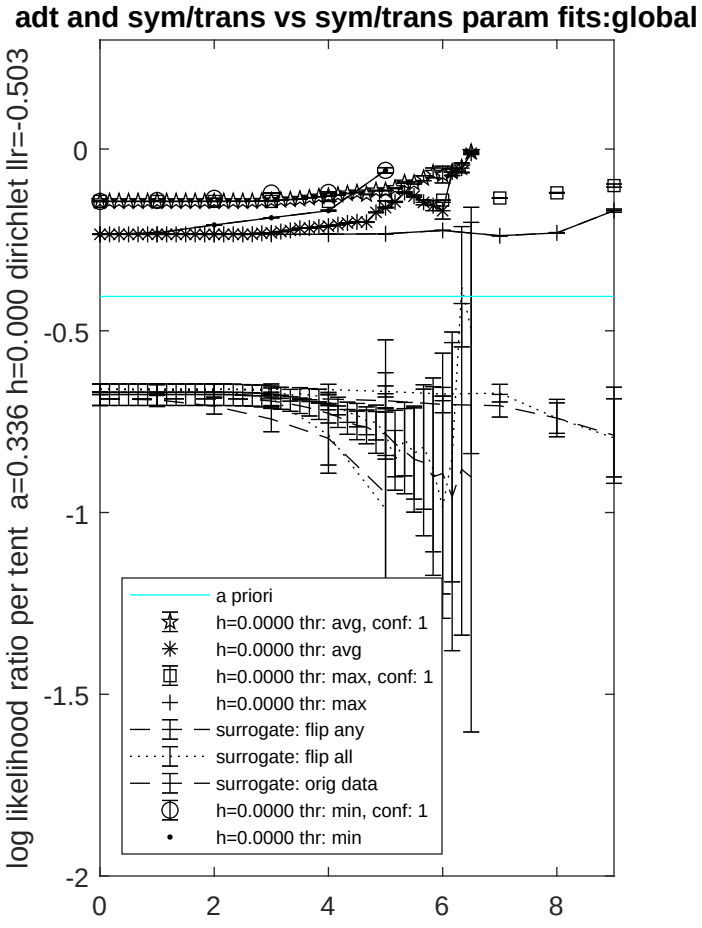
./psg_data/bc6pt_choices_MO_sess01_10 eb: 1 SD, fitted h sel_bxrand



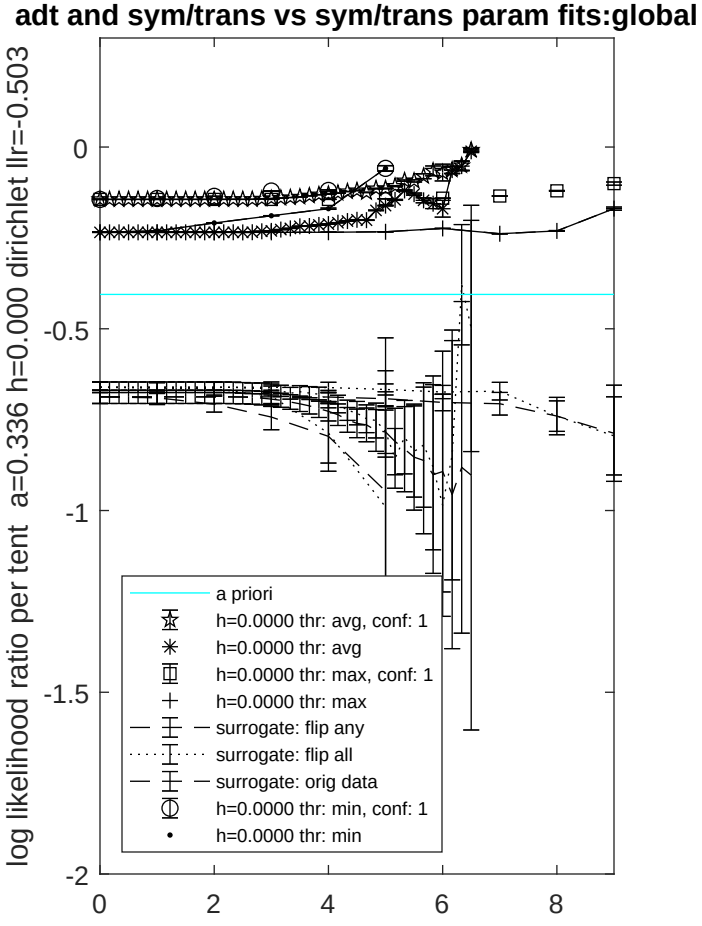




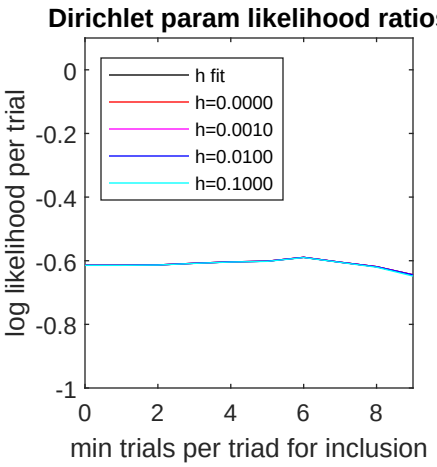
./psg_data/bc6pt_choices_MC_sess01_10 eb: 1 SD sel_cXrand

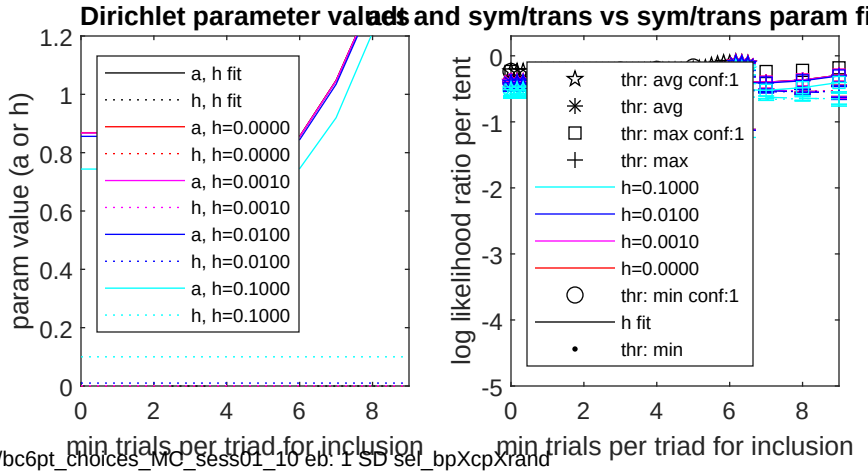


./psg_data/bc6pt_choices_MO_sess01_1e eb: 1 SD, fixed h set_cXrand

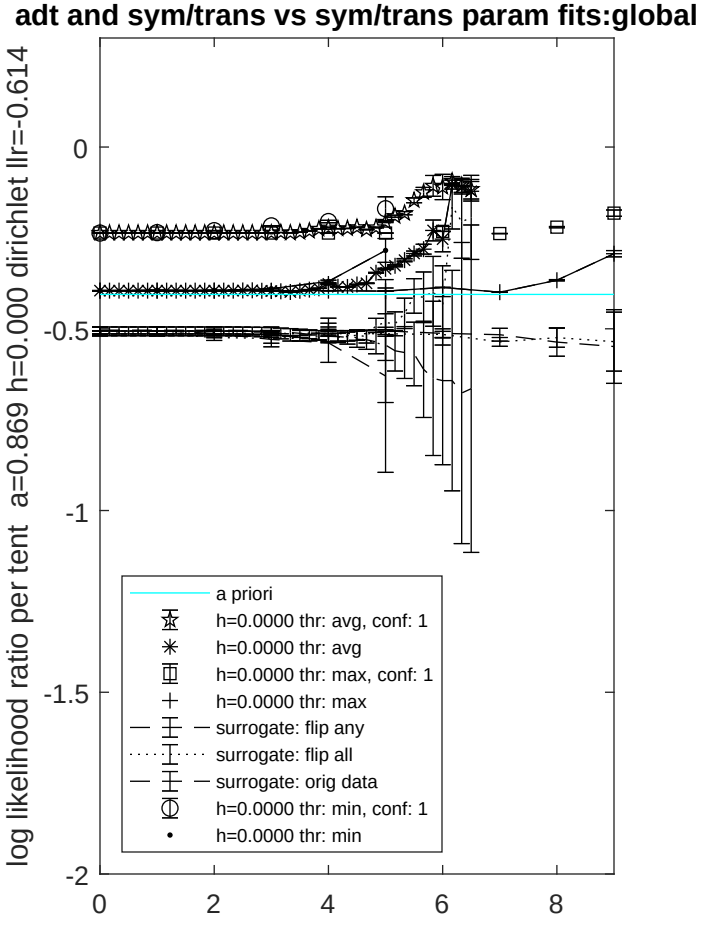


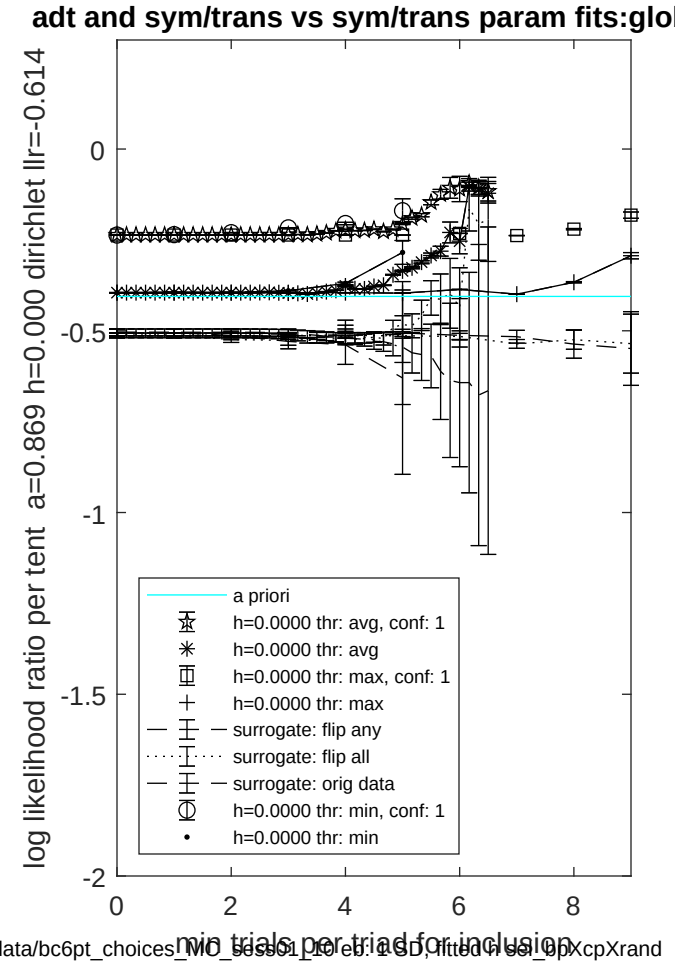
./psg_data/bc6pt_choices_MO_sess01_10 eb: 1 SD, fitted h sel_cXrand

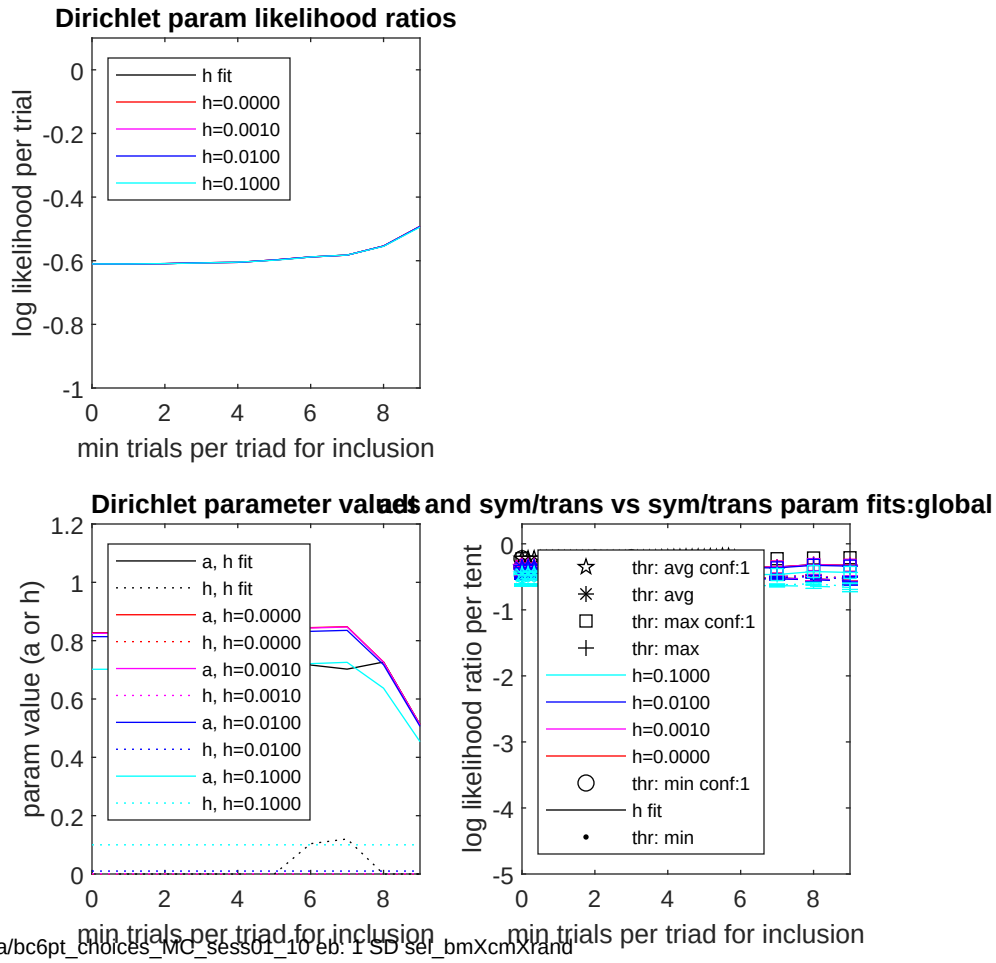




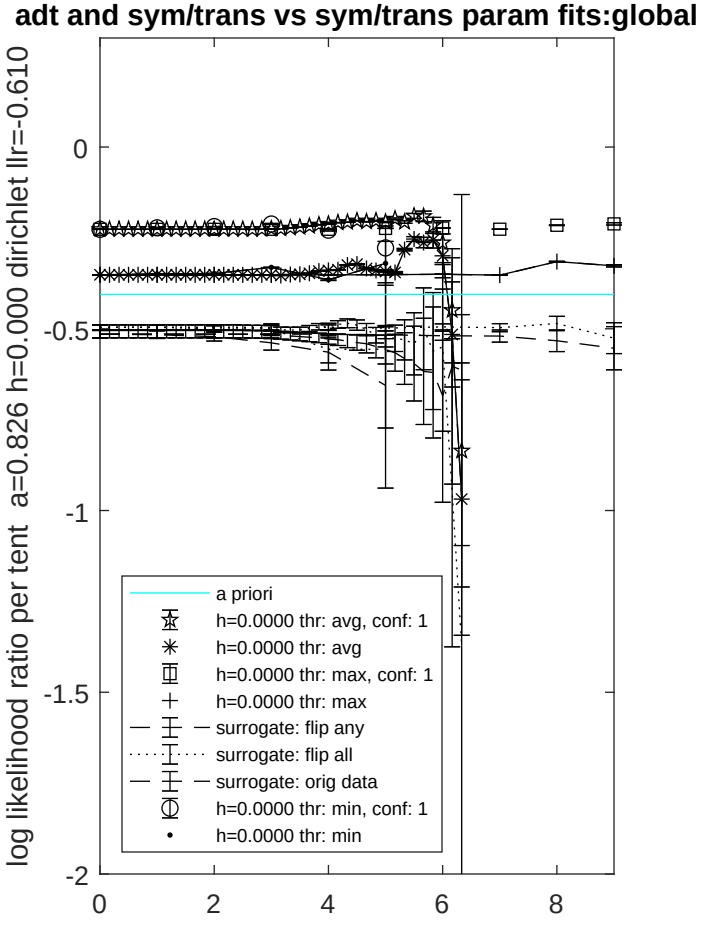
./psg_data/bc6pt_choices_MC_sess01_10 eb: 1 SD sel_bpXcpXrand

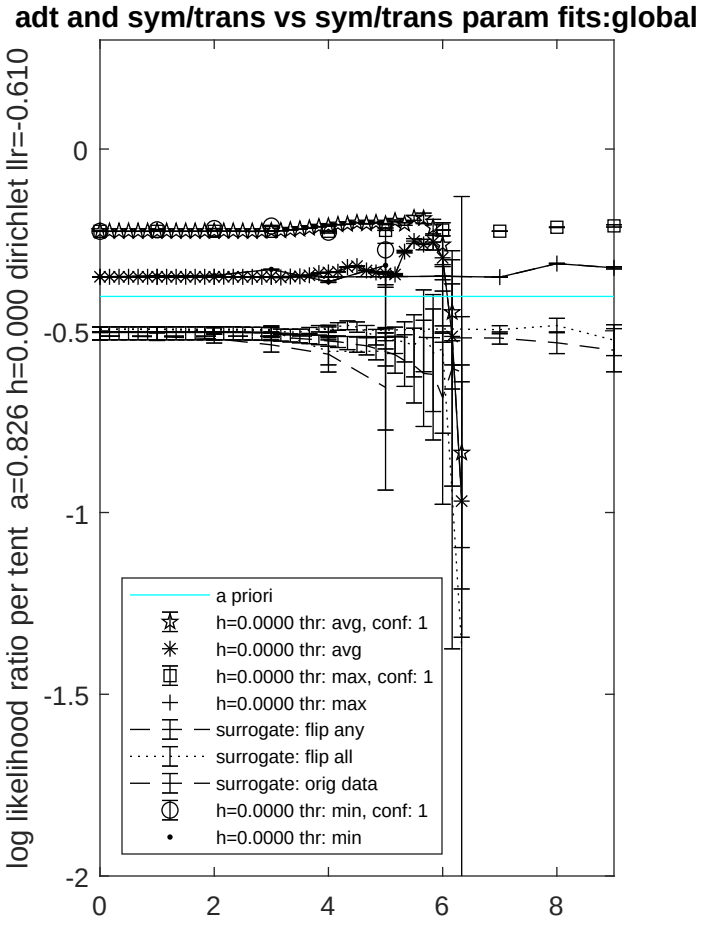

adt and sym/trans vs sym/trans param fits:global
log likelihood ratio per tent a=0.869 h=0.000 dirichlet llr=-0.614
0
-0.5
-1
-1.5
-2
0
2
4
6
8
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
./psg_data/bc6pt_choices_MO_sess01_10 eb: 1 SD, fixed h ser_bpXcpXrand

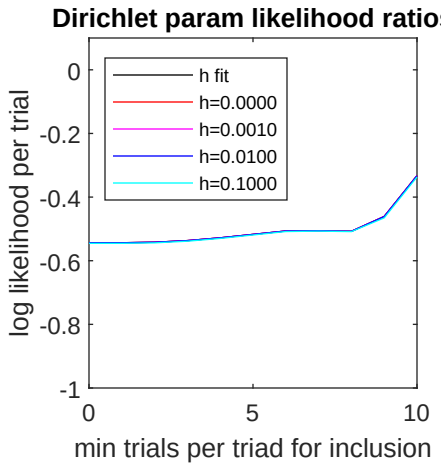

adt and sym/trans vs sym/trans param fits:global
log likelihood ratio per tent a=0.869 h=0.000 dirichlet llr=-0.614
0
-0.5
-1
-1.5
-2
0
2
4
6
8
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
./psg_data/bc6pt_choices_MO_sess01_10 eb: 1 SD, fitted h sel_bpXcpXrand

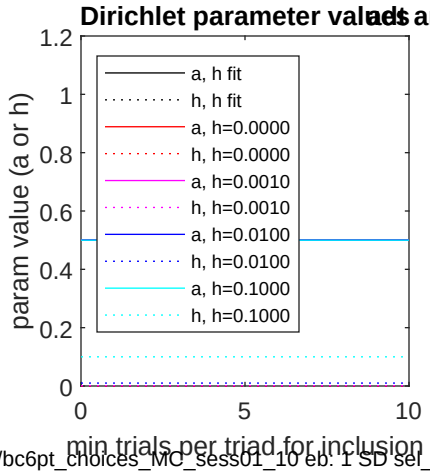


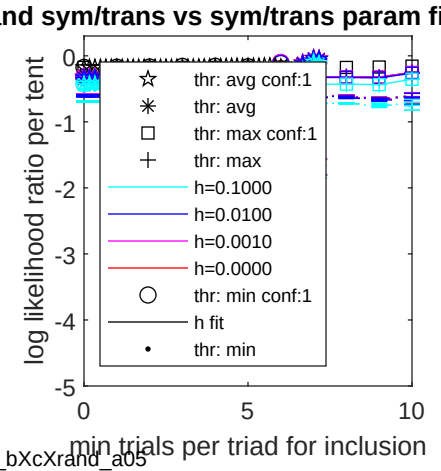
./psg_data/bc6pt_choices_MC_sess01_10 eb: 1 SD sel_bmXcmXrand


adt and sym/trans vs sym/trans param fits:global
log likelihood ratio per tent a=0.826 h=0.000 dirichlet llr=-0.610
0
-0.5
-1
-1.5
-2
0
2
4
6
8
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
./psg_data/bc6pt_choices_MO_sess01_10 eb: 1 SD, fixed h set_bmXcmXrand

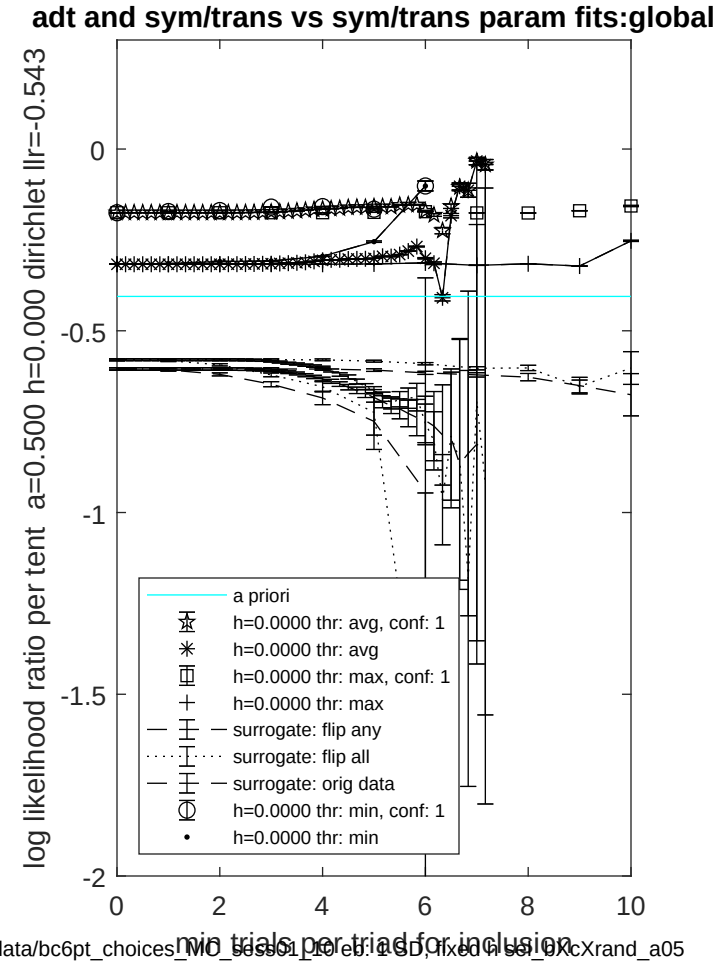

adt and sym/trans vs sym/trans param fits:global
log likelihood ratio per tent a=0.826 h=0.000 dirichlet llr=-0.610
0
-0.5
-1
-1.5
-2
0
2
4
6
8
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
./psg_data/bc6pt_choices_MO_sess01_10 eb: 1 SD, fitted h sel_bmXcmXrand



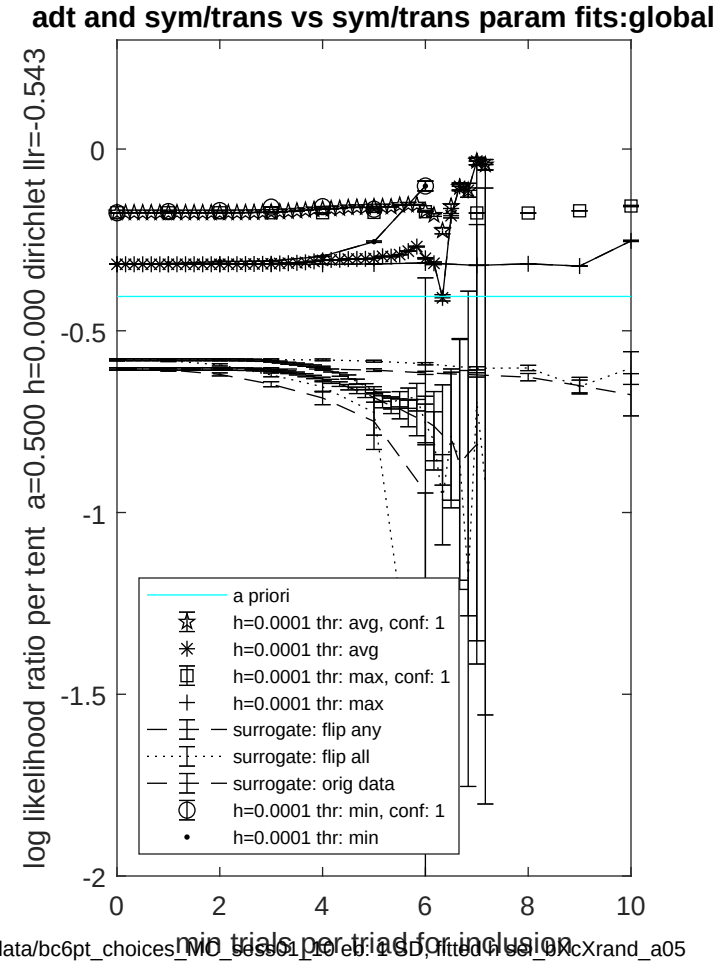




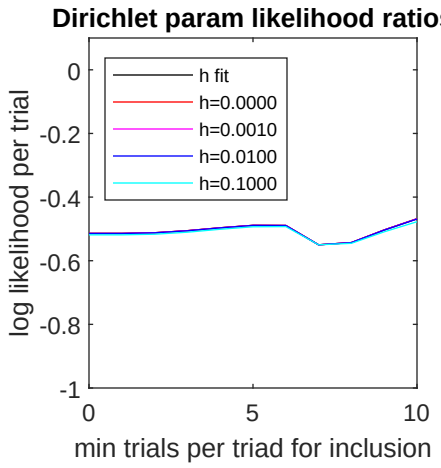
./psg_data/bc6pt_choices_MC_sess01_10 eb: 1 SD sel_bXcXrand_a05

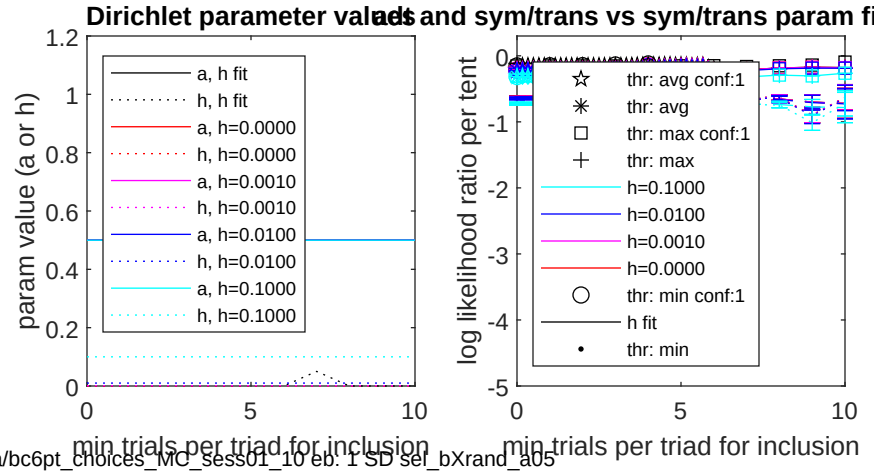


./psg_data/bc6pt_choices_MC_sess01_10 eb: 1 SD, fixed h sel_bXcXrand_a05



./psg_data/bc6pt_choices_MO_sess01_10 eb: 1 SD, fitted h sel_bXcXrand_a05





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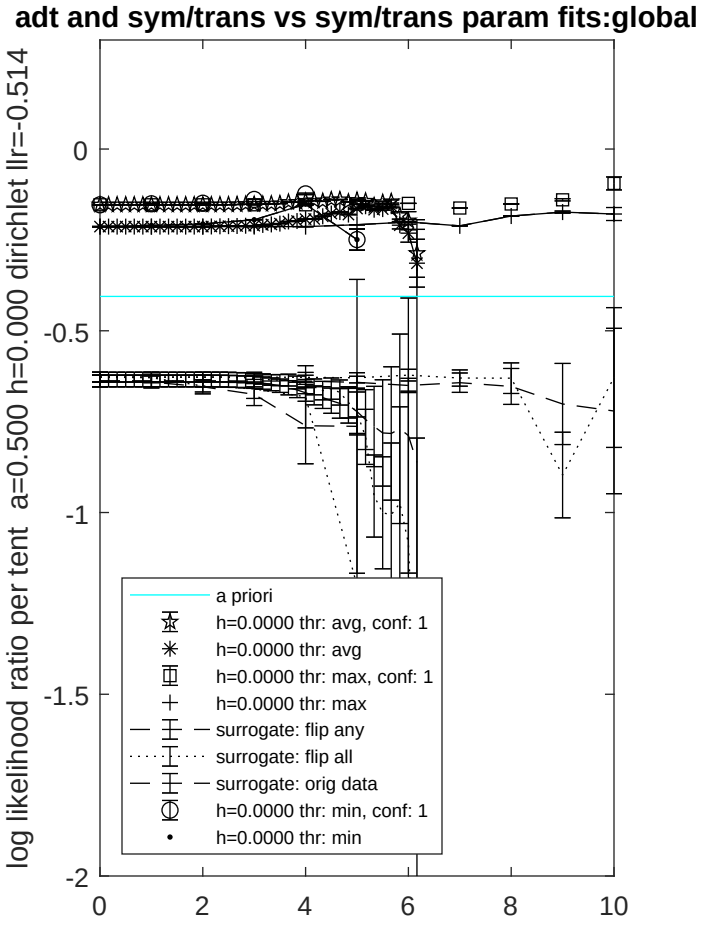
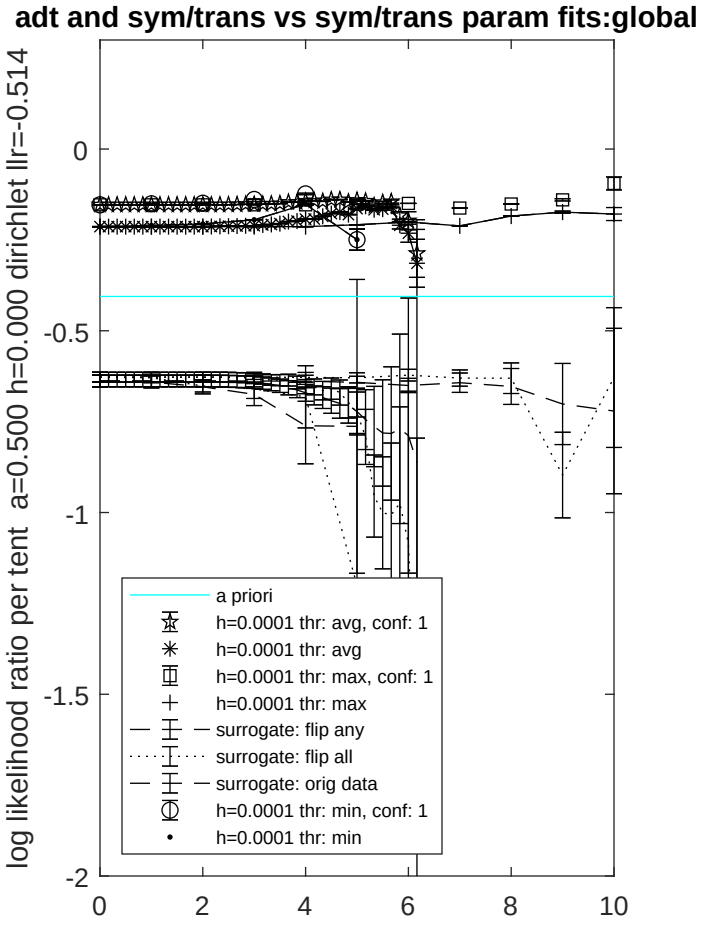
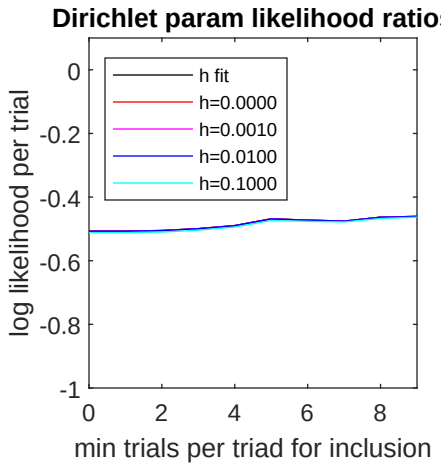
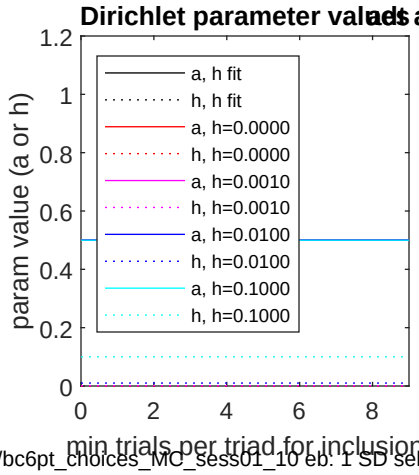


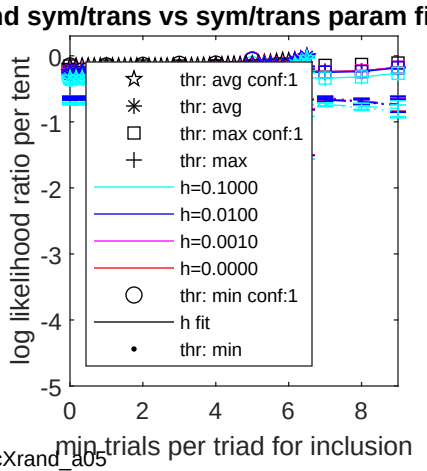
./psg_data/bc6pt_choices_MC_sess01_10 eb: 1 SD, fixed h set_bXrand_a05



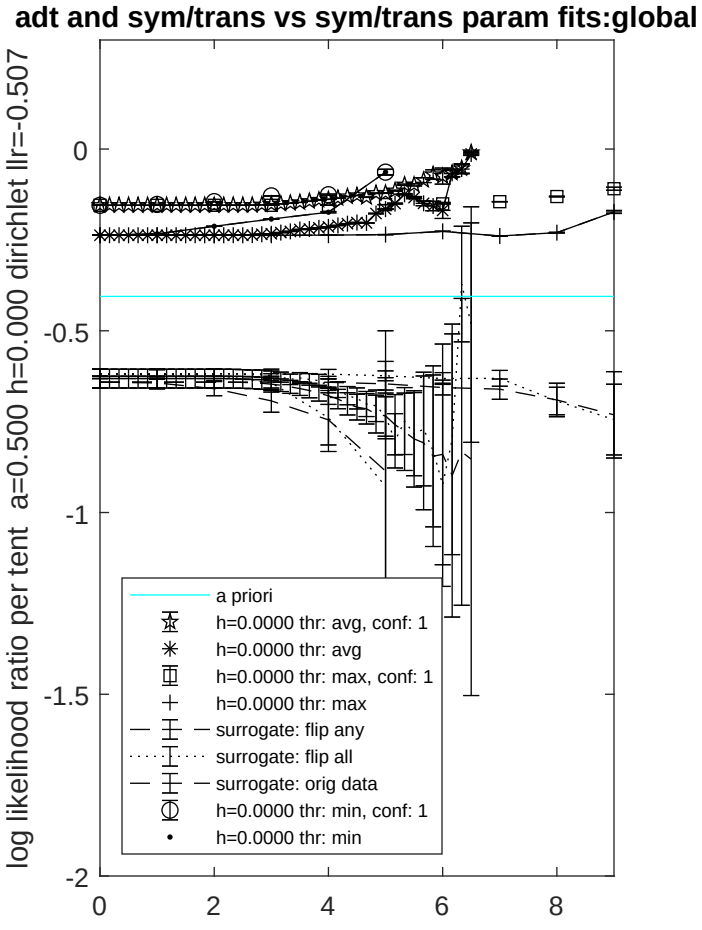
./psg_data/bc6pt_choices_MO_sess01_10 eb: 1 SD, fitted h sel_bXrand_a05

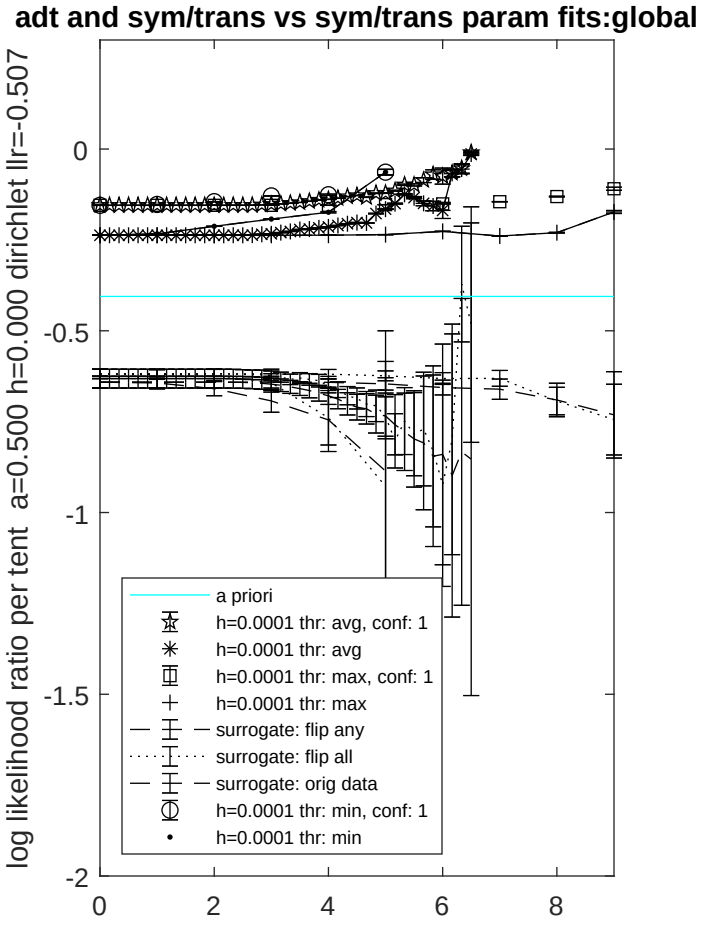






./psg_data/bc6pt_choices_MC_sess01_10 eb: 1 SD sel_cXrand_a05


adt and sym/trans vs sym/trans param fits:global
log likelihood ratio per tent a=0.500 h=0.000 dirichlet llr=-0.507
0
-0.5
-1
-1.5
-2
0
2
4
6
8
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
./psg_data/bc6pt_choices_MC_sess01_1c eb: 1 SD, fixed h set_cXrand_a05



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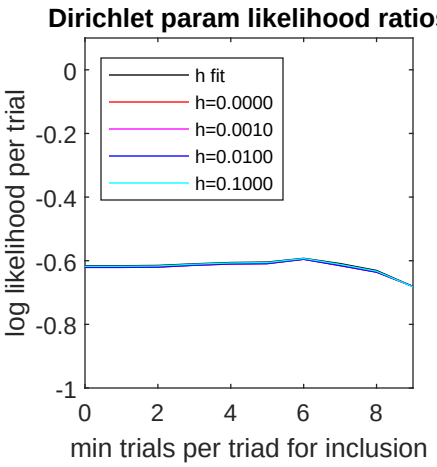
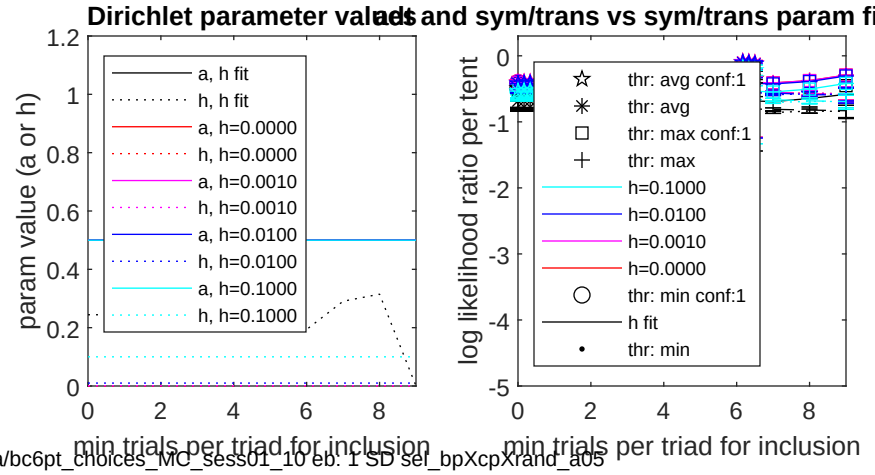
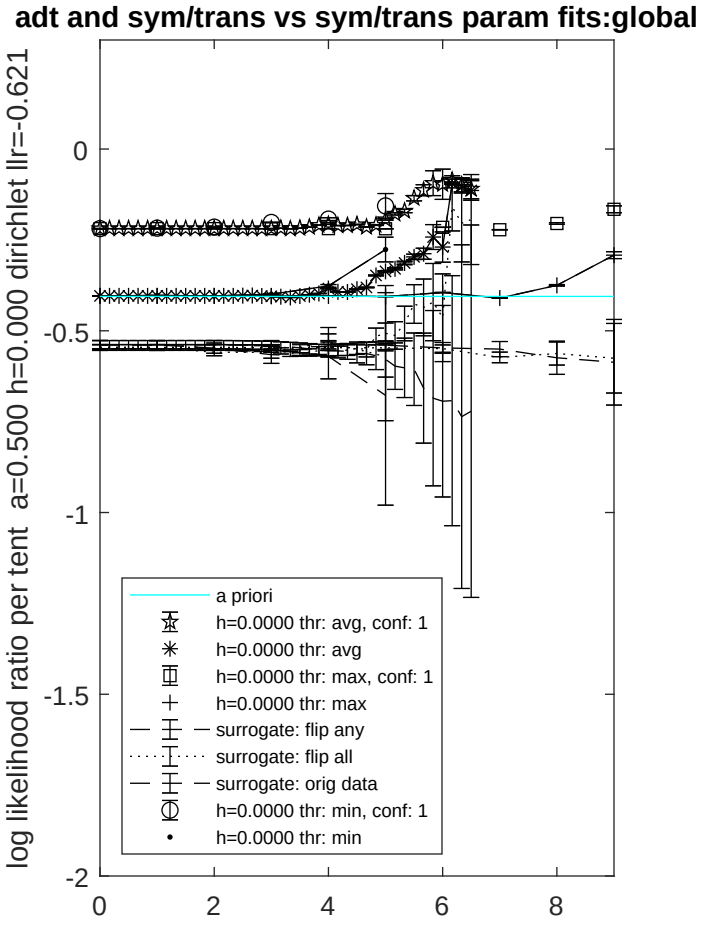
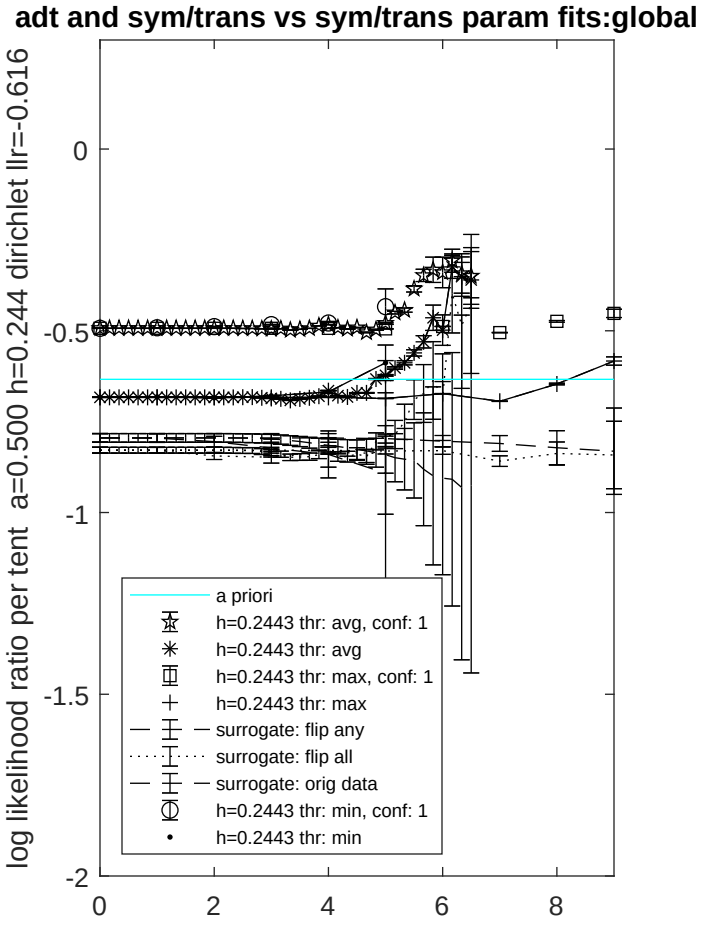


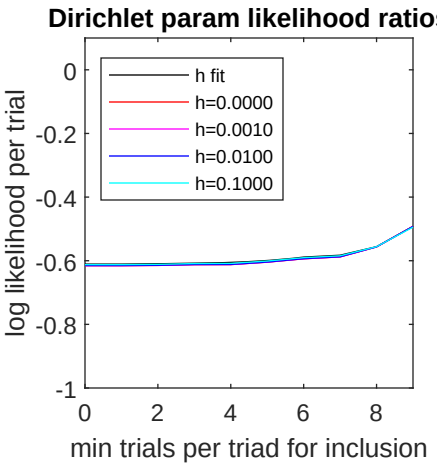


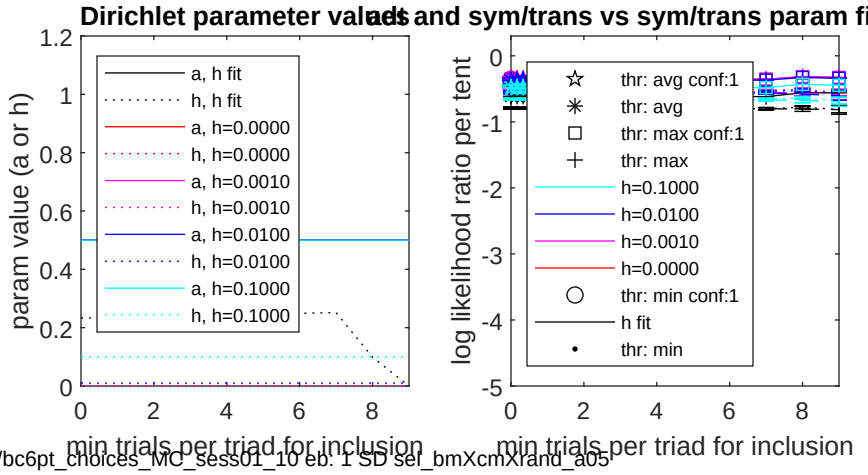
./psg_data/bc6pt_choices_MC_sess01_10 eb: 1 SD sel_bpXcpXrand_a05



./psg_data/bc6pt_choices_MO_sess01_to_eb_1SD_fixed_ser_bpXcpXrand_a05


adt and sym/trans vs sym/trans param fits:global
log likelihood ratio per tent a=0.500 h=0.244 dirichlet llr=-0.616
0
-0.5
-1
-1.5
-2
0
2
4
6
8
min trials per triad for inclusion
a priori
h=0.2443 thr: avg, conf: 1
h=0.2443 thr: avg
h=0.2443 thr: max, conf: 1
h=0.2443 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.2443 thr: min, conf: 1
h=0.2443 thr: min
./psg_data/bc6pt_choices_MO_sess01_10 eb: 1 SD, fitted h sel_bpXcpXrand_a05





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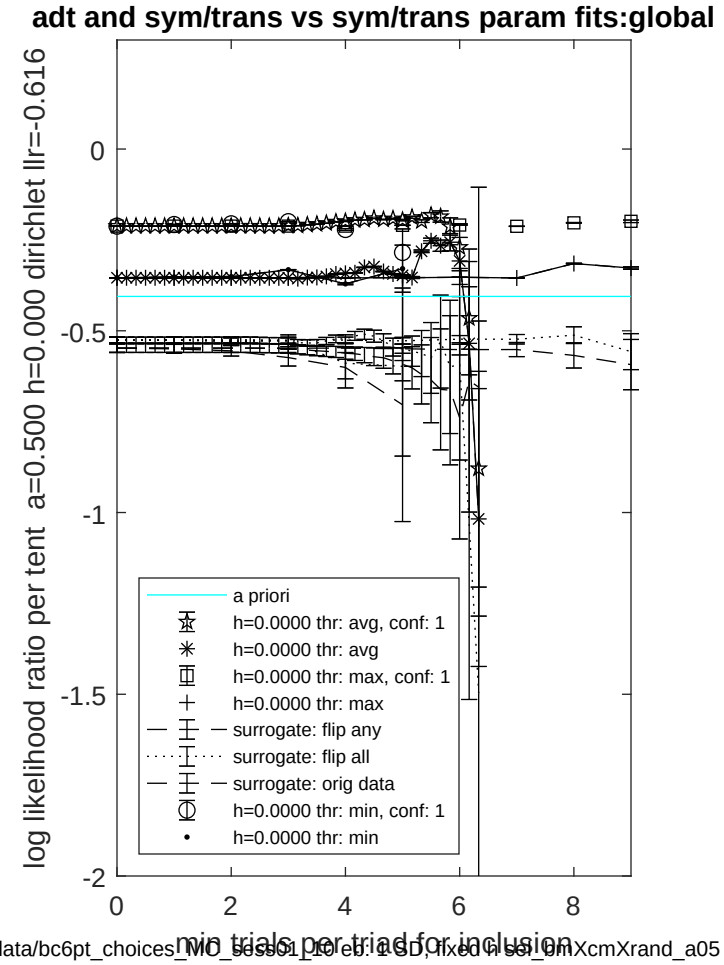
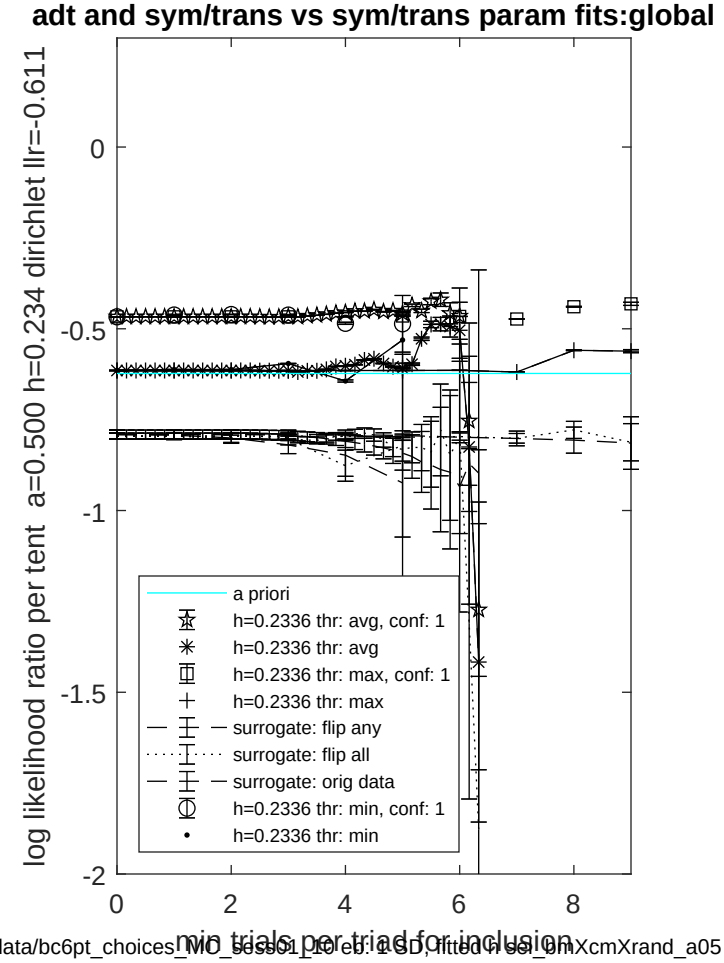
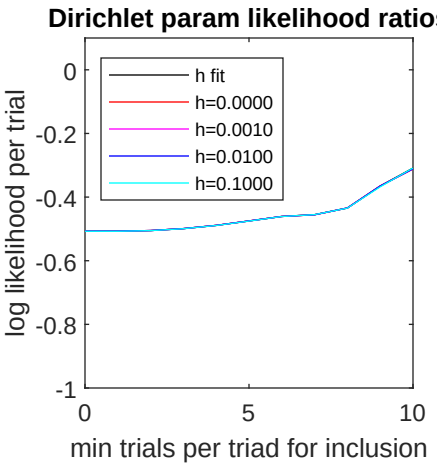
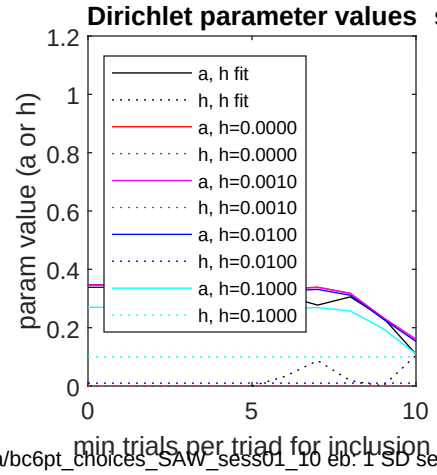
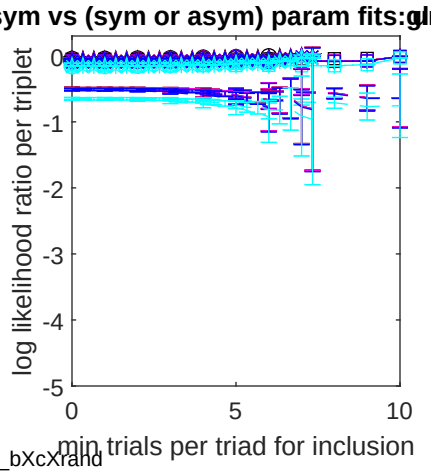
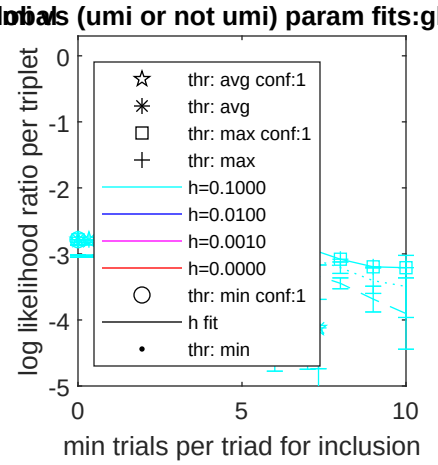

adt and sym/trans vs sym/trans param fits:global
log likelihood ratio per tent a=0.500 h=0.000 dirichlet llr=-0.616
0
-0.5
-1
-1.5
-2
0
2
4
6
8
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
./psg_data/bc6pt_choices_MO_sess01_1c eb: 1 SD, fixed h set_bmXcmXrand_a05


adt and sym/trans vs sym/trans param fits:global
log likelihood ratio per tent a=0.500 h=0.234 dirichlet llr=-0.611
0
-0.5
-1
-1.5
-2
0
2
4
6
8
min trials per triad for inclusion
a priori
h=0.2336 thr: avg, conf: 1
h=0.2336 thr: avg
h=0.2336 thr: max, conf: 1
h=0.2336 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.2336 thr: min, conf: 1
h=0.2336 thr: min
./psg_data/bc6pt_choices_MO_sess01_10 eb: 1 SD, fitted h sel_bmXcmXrand_a05

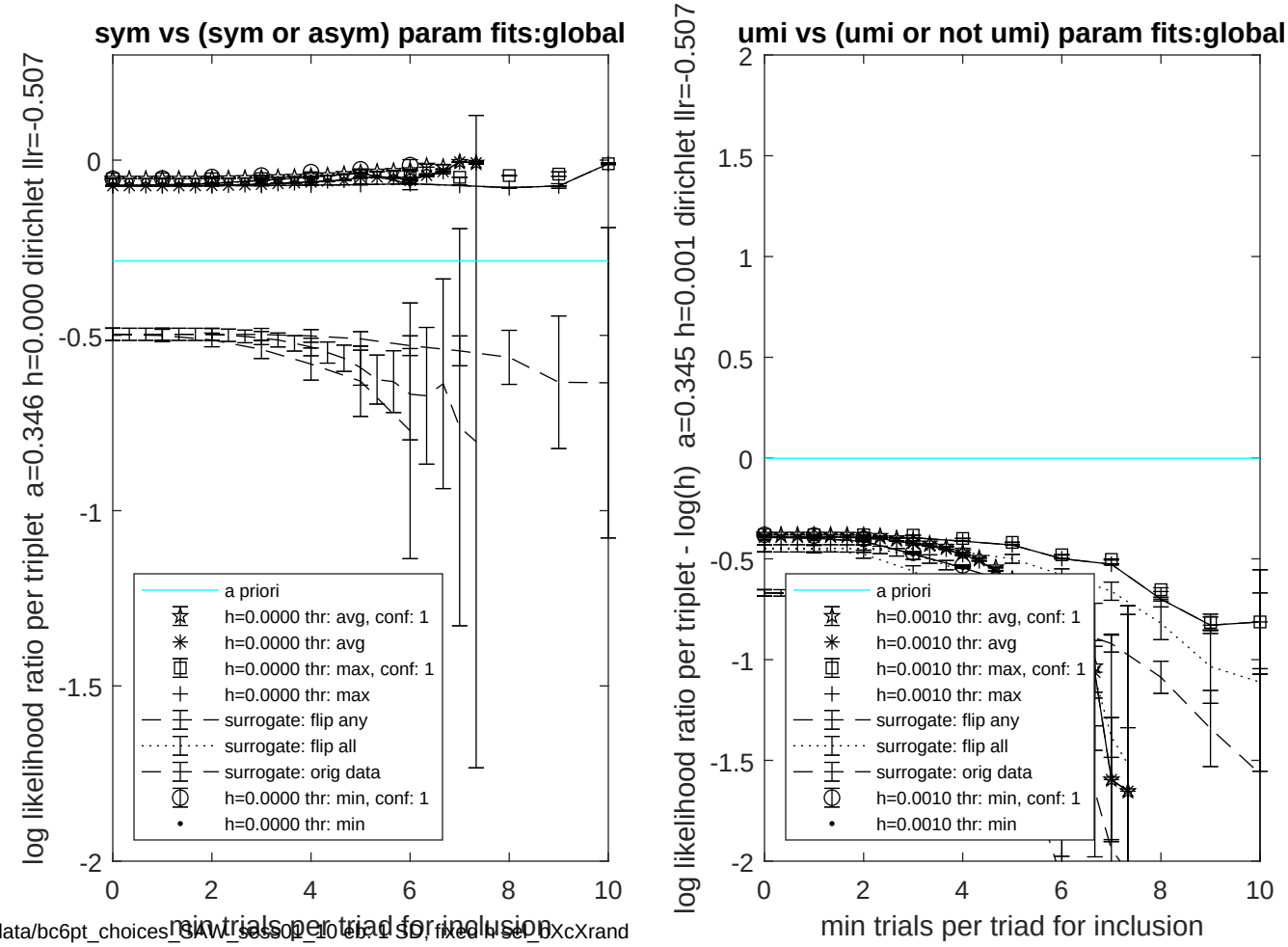




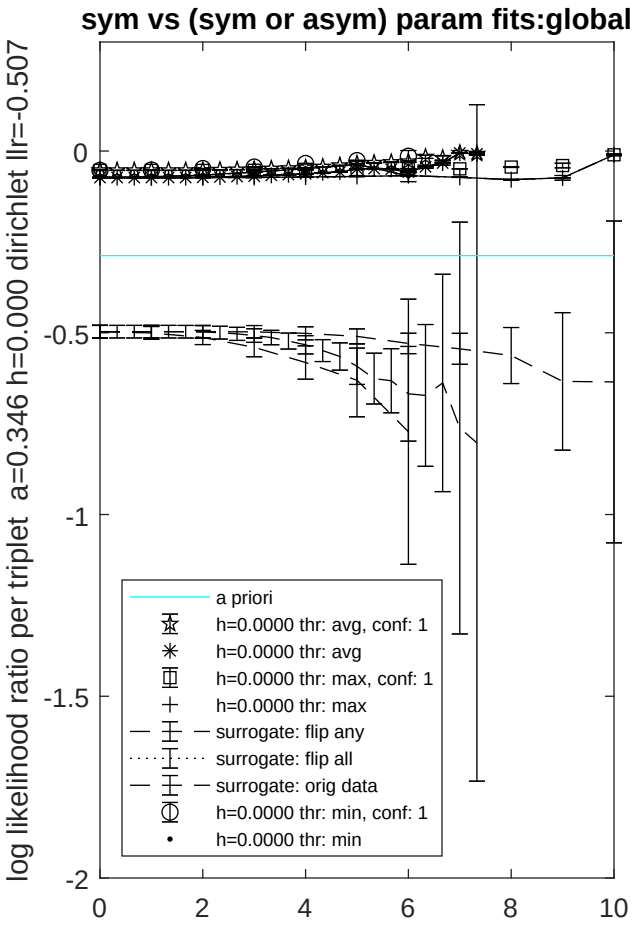


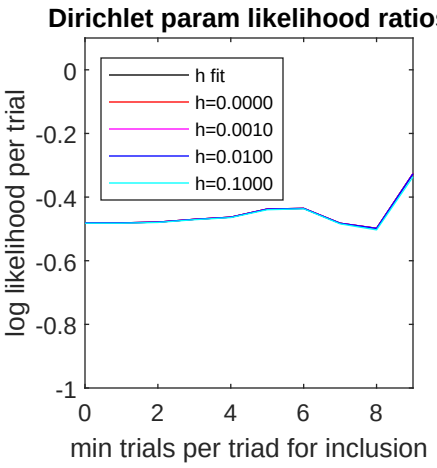


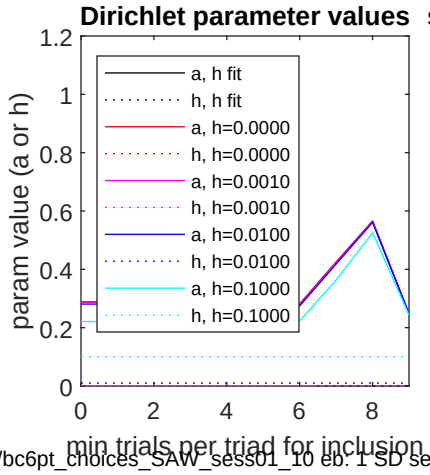
./psg_data/bc6pt_choices_SAW_sess01_10 eb: 1 SD sel_bXcXrand

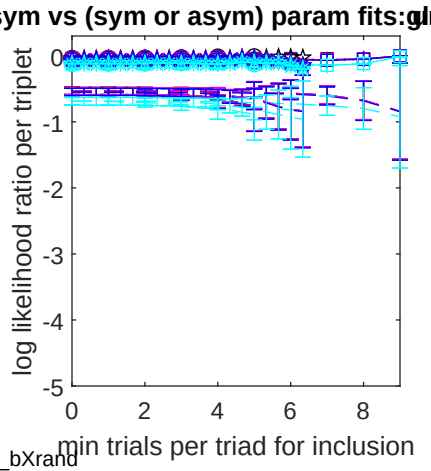


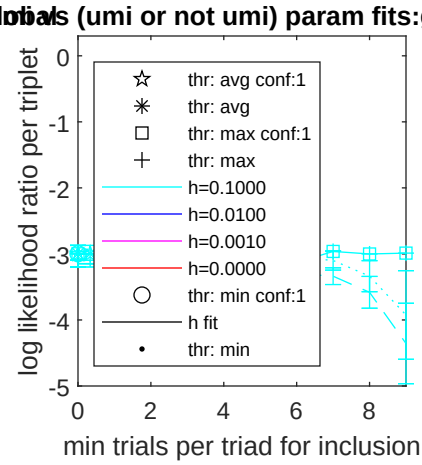
./psg_data/bc6pt_choices_SAW_sess01_10 eb: 1 SD, fixed h sel_bXcXrand


sym vs (sym or asym) param fits:global
log likelihood ratio per triplet a=0.346 h=0.000 dirichlet llr=-0.507
0
-0.5
-1
-1.5
-2
0
2
4
6
8
10
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
./psg_data/bc6pt_choices_SAW_sess0p_10 eb: 1 SD, fitted h sel_bXcXrand

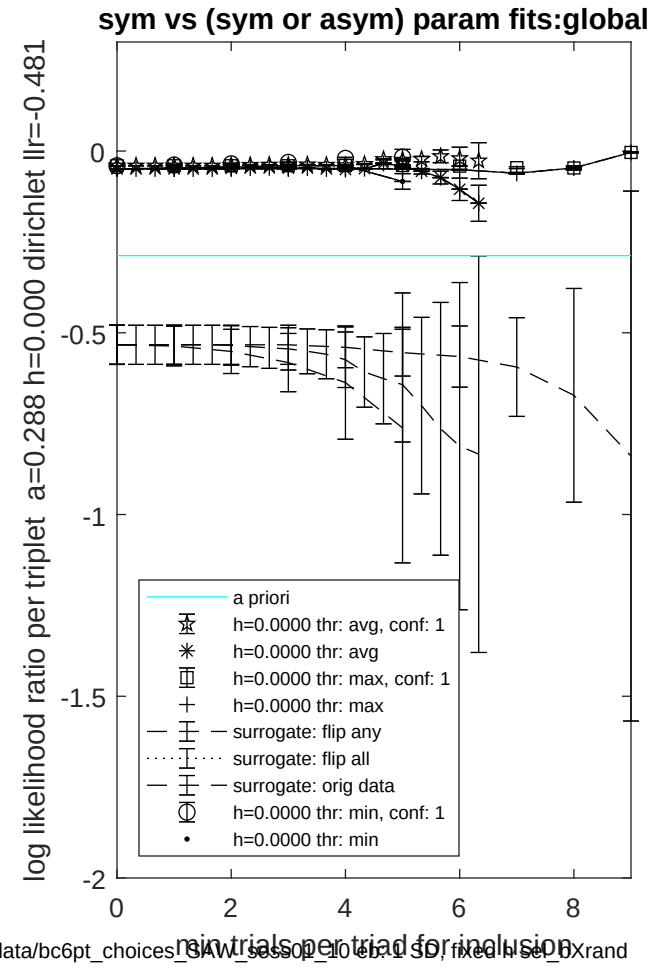


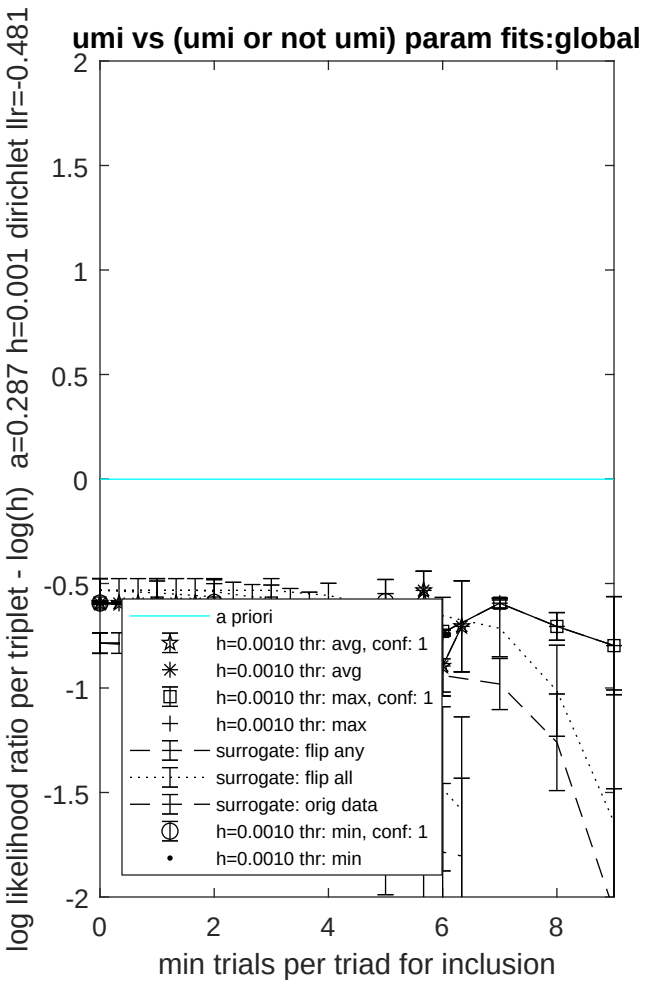




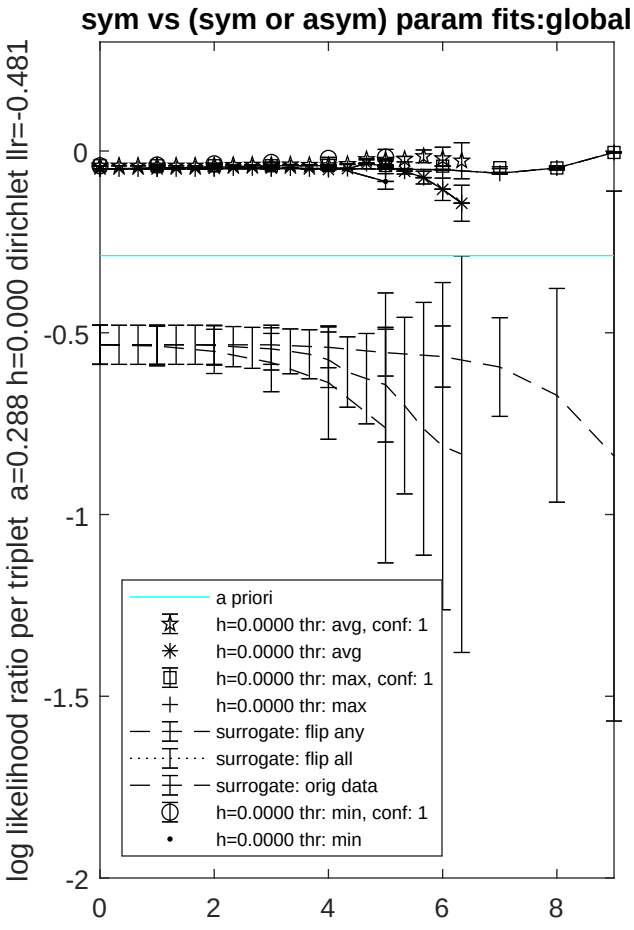


./psg_data/bc6pt_choices_SAW_sess01_10 eb: 1 SD sel_bXrand

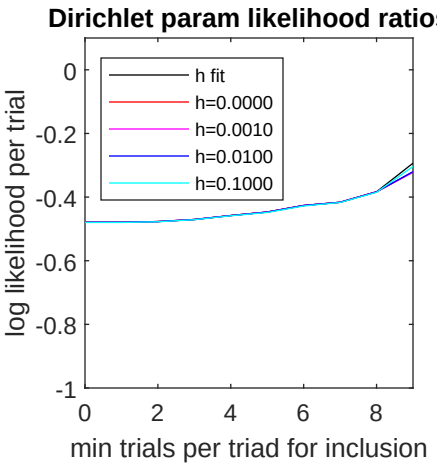


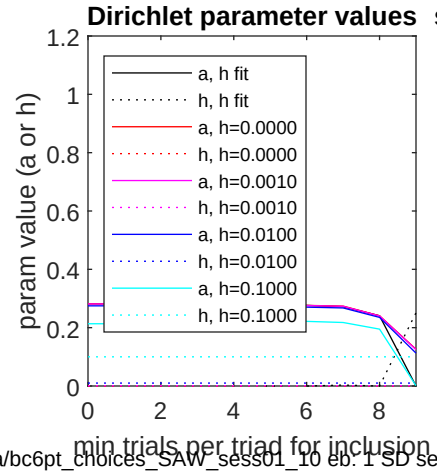


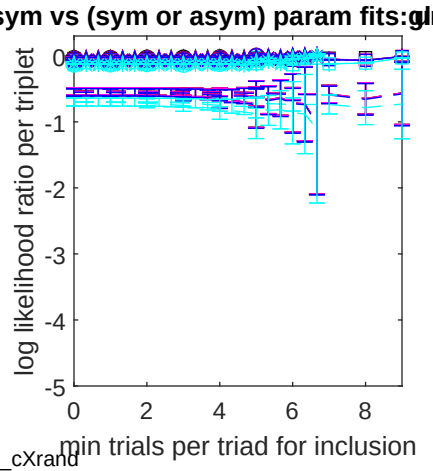
./psg_data/bc6pt_choices_SAW_sess0[_10 eb.1 SD, fixed h set_0Xrand

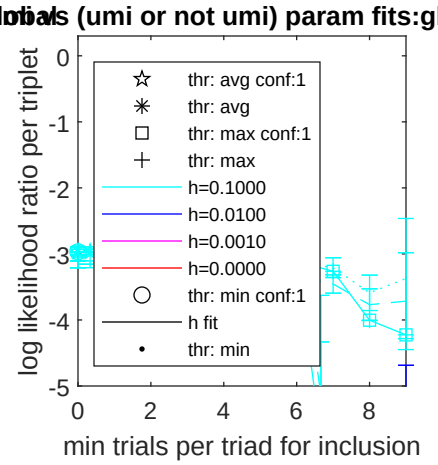


./psg_data/bc6pt_choices_SAW_sess01_10 eb:1 SD, fitted h set_bXrand

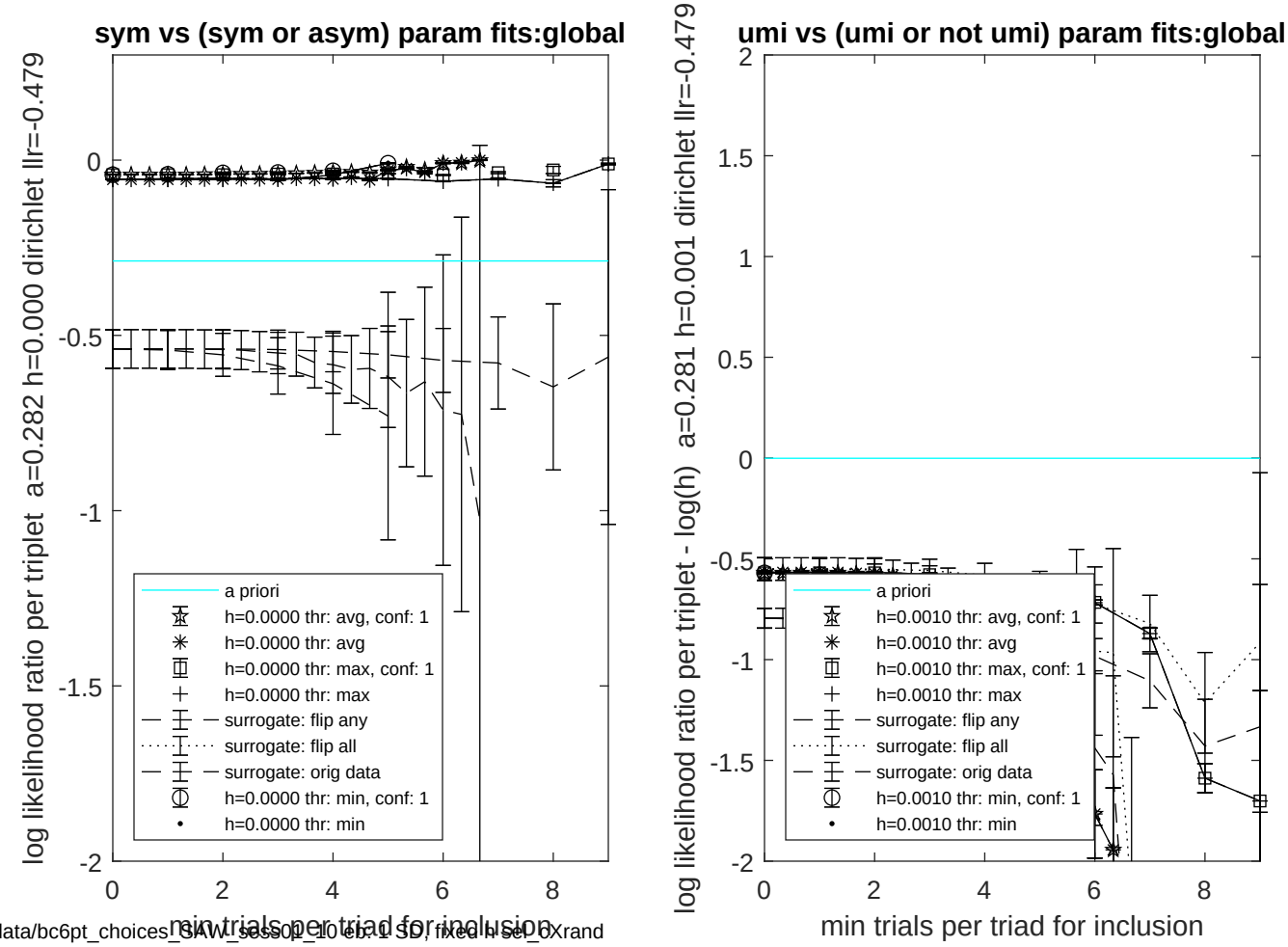




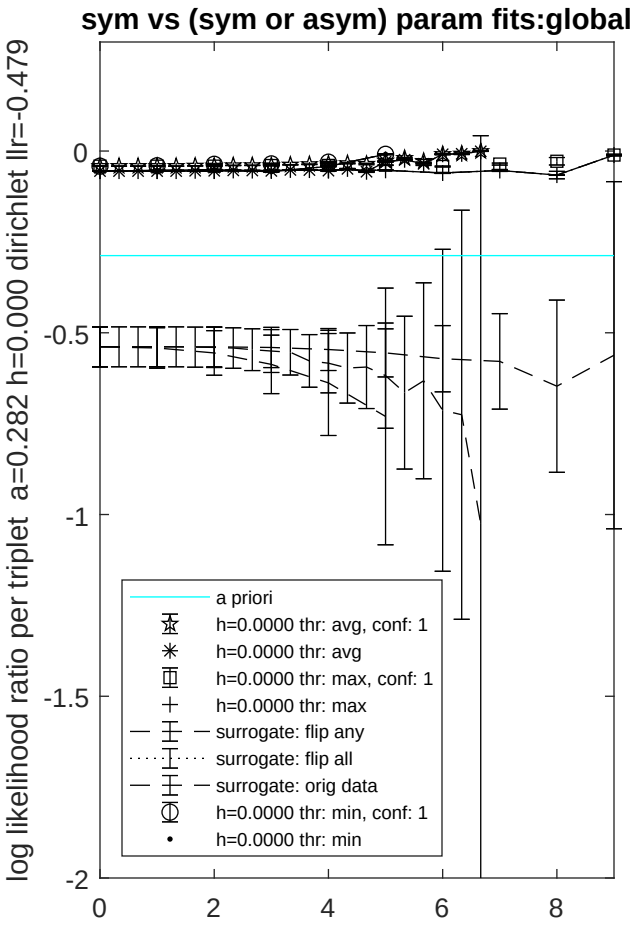




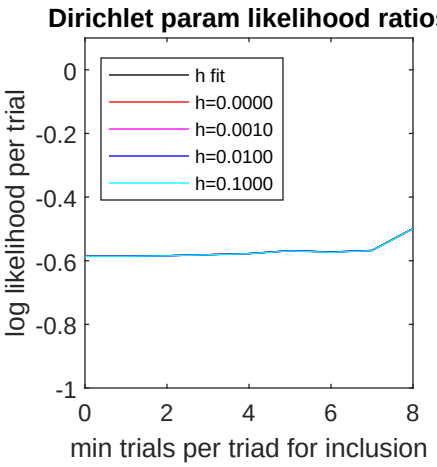
./psg_data/bc6pt_choices_SAW_sess01_10 eb: 1 SD sel_cXrand

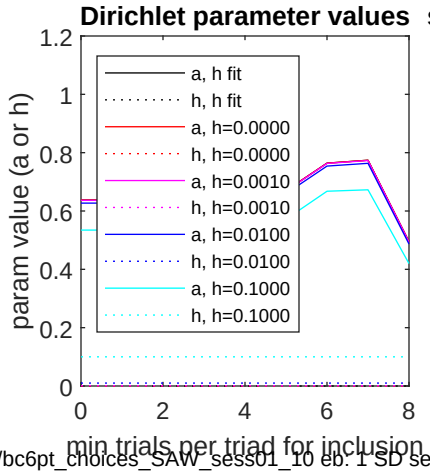


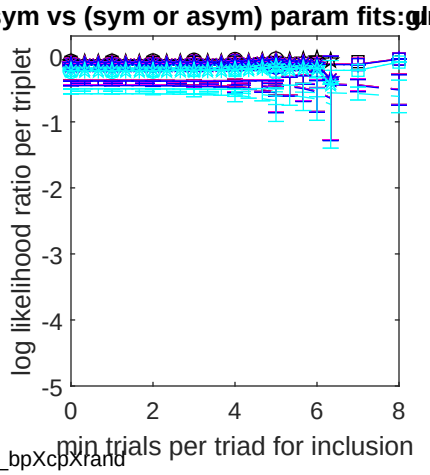
./psg_data/bc6pt_choices_SAW_sess01_10 eb: 1 SD, fixed h sel_0Xrand

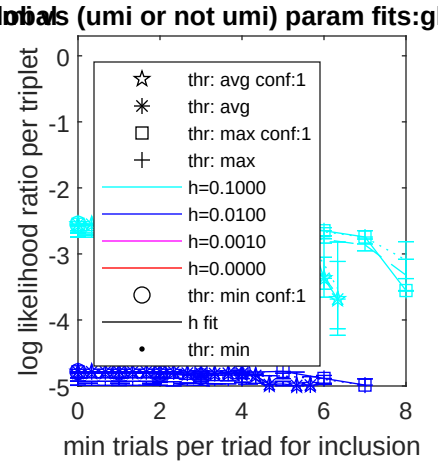


./psg_data/bc6pt_choices_SAW_sess01_10 eb:1 SD, fitted h set_0Xrand









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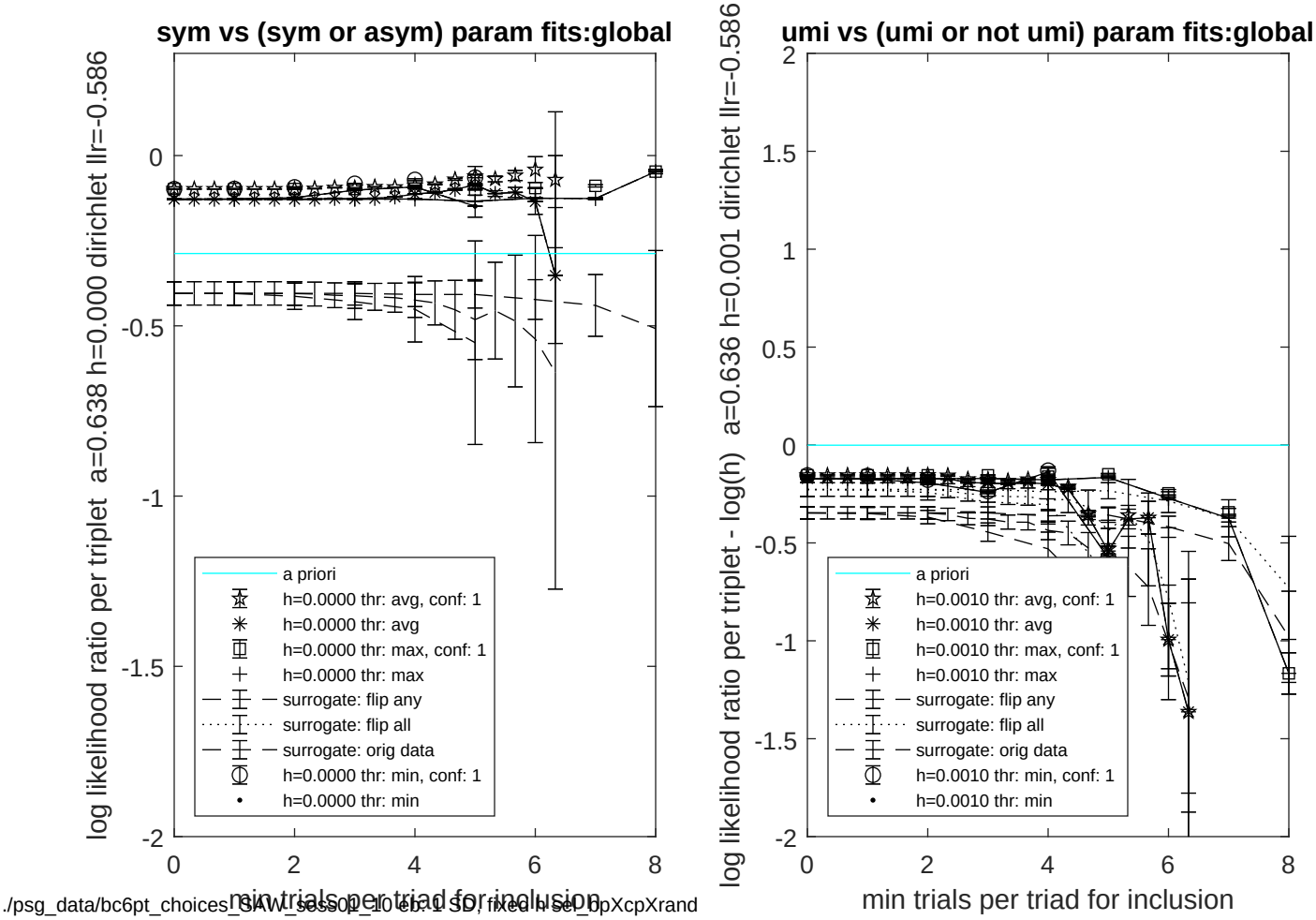
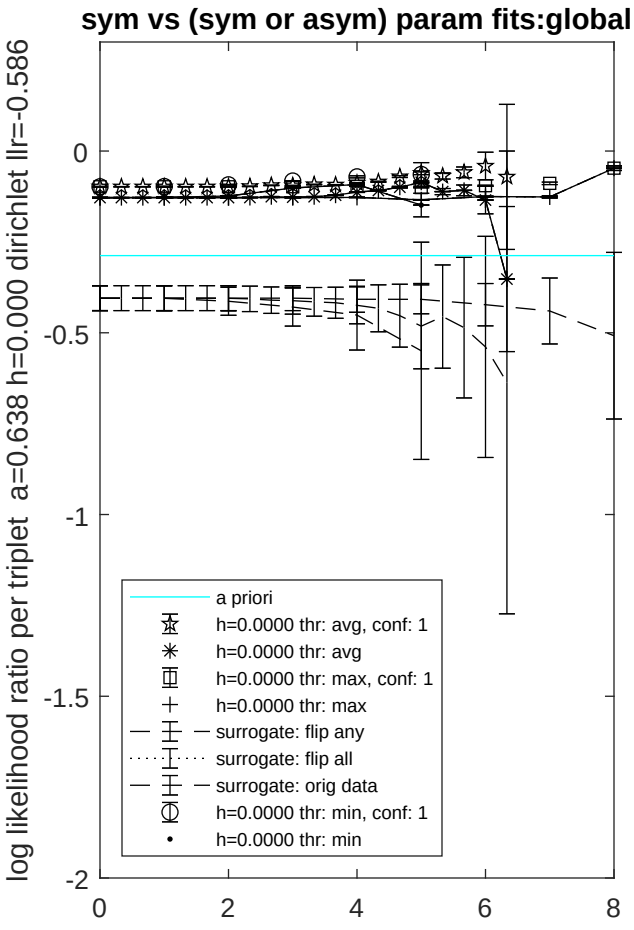

sym vs (sym or asym) param fits:global
log likelihood ratio per triplet a=0.638 h=0.000 dirichlet llr=-0.586
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
./psg_data/bc6pt_choices_SAW_sess01_10 eb: 1 SD, fixed h sel_bpXcpXrand
umi vs (umi or not umi) param fits:global
log likelihood ratio per triplet - log(h) a=0.636 h=0.001 dirichlet llr=-0.586
min trials per triad for inclusion
a priori
h=0.0010 thr: avg, conf: 1
h=0.0010 thr: avg
h=0.0010 thr: max, conf: 1
h=0.0010 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0010 thr: min, conf: 1
h=0.0010 thr: min


sym vs (sym or asym) param fits:global
log likelihood ratio per triplet a=0.638 h=0.000 dirichlet llr=-0.586
0
-0.5
-1
-1.5
-2
0
2
4
6
8
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
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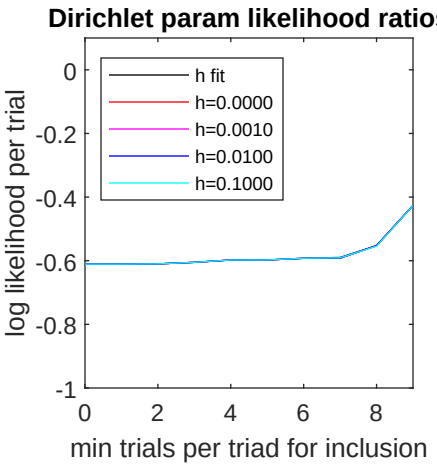

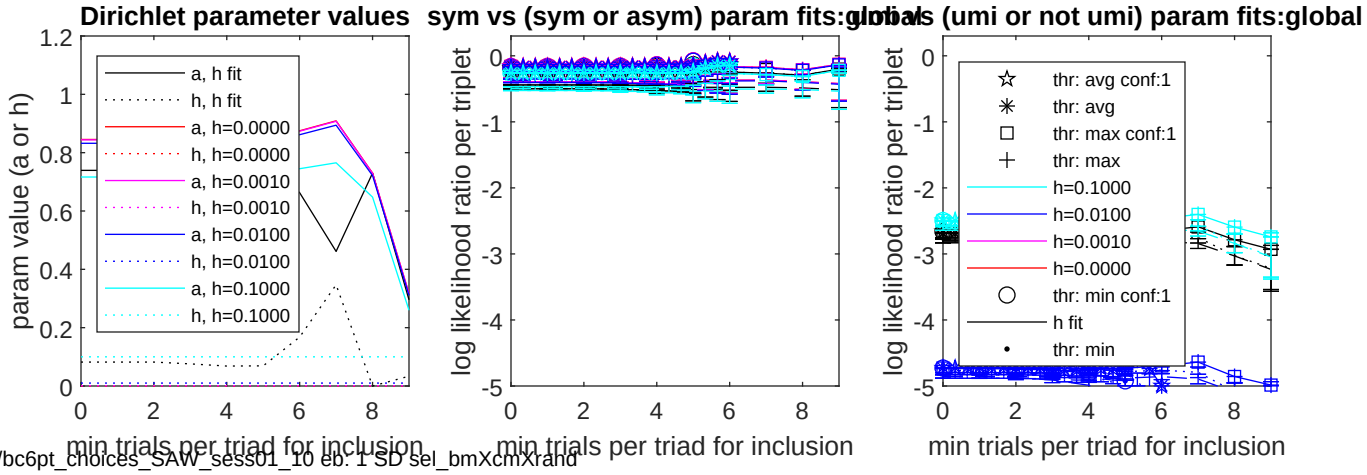

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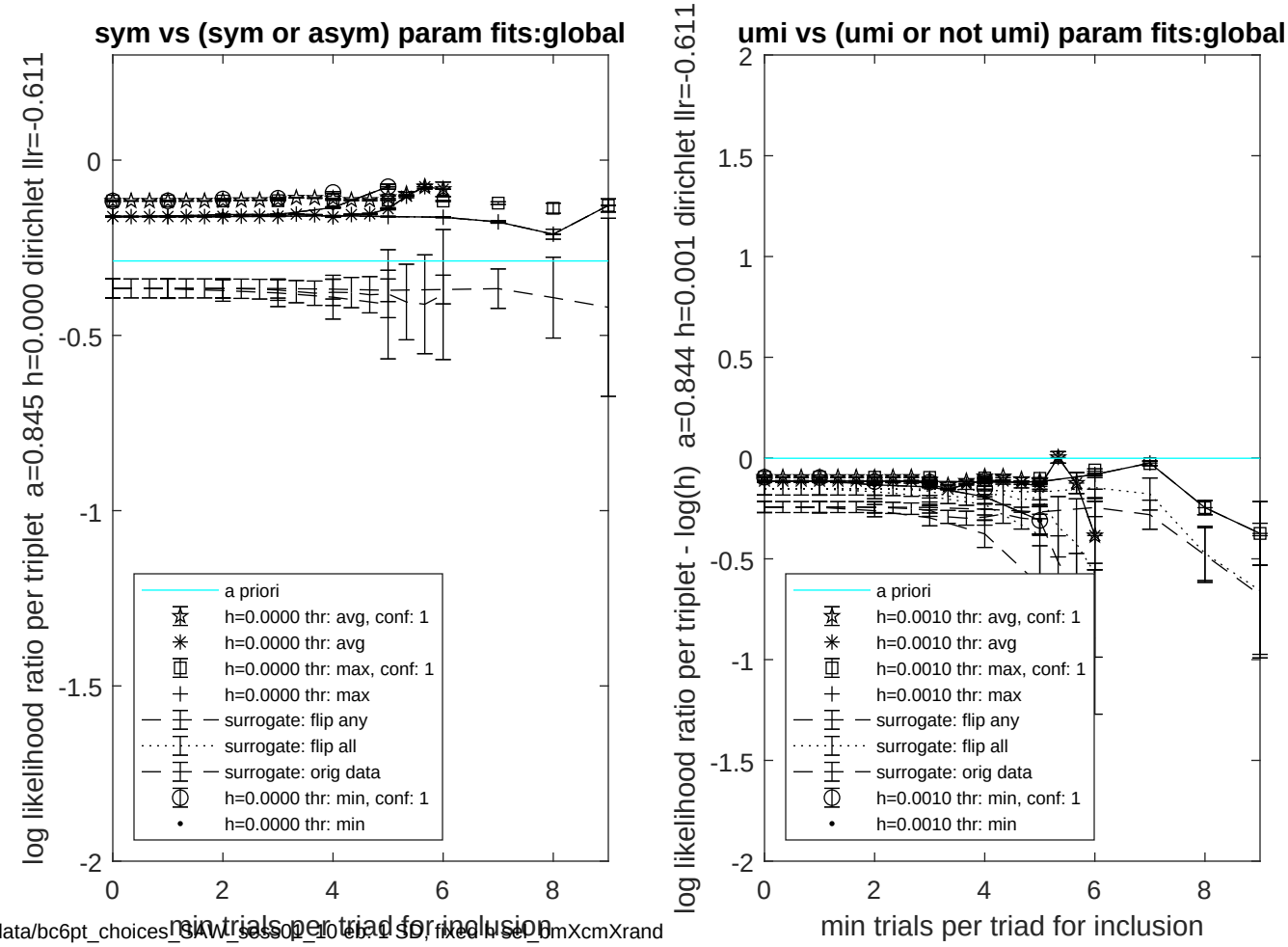

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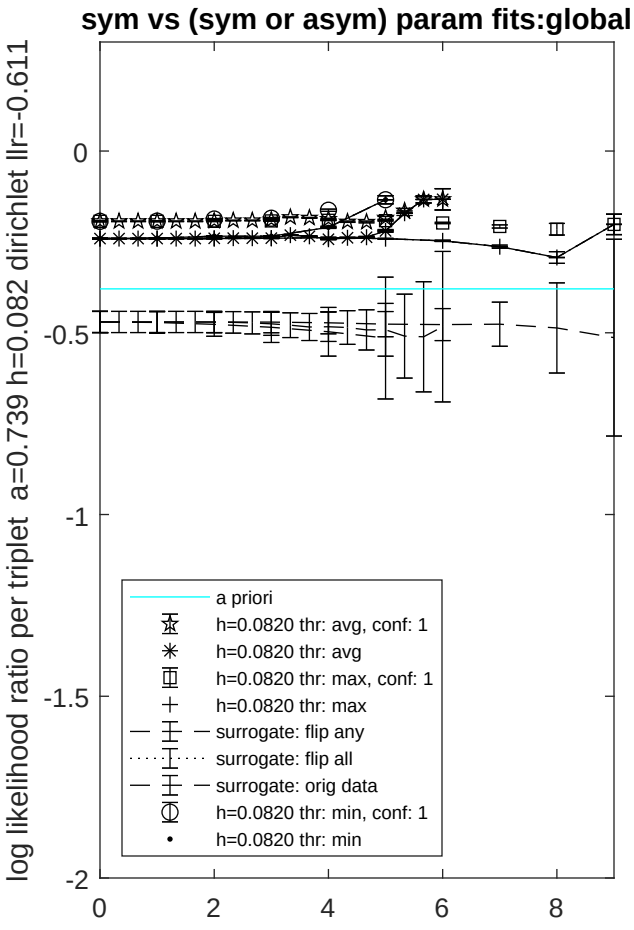

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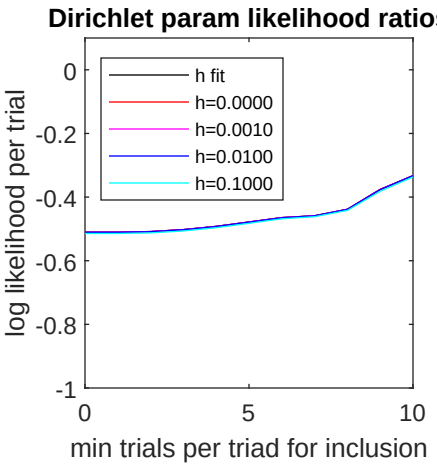

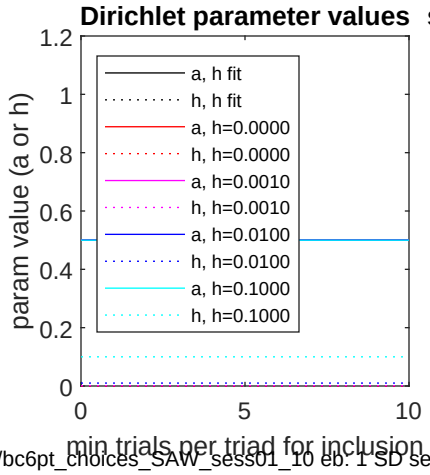

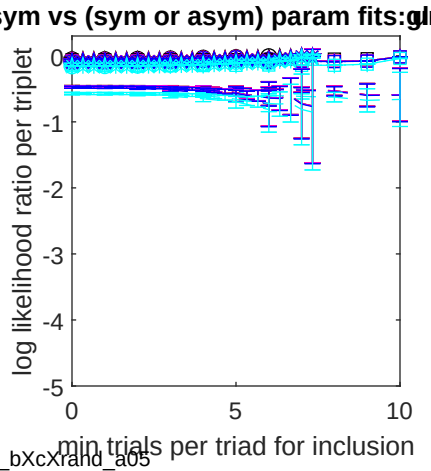

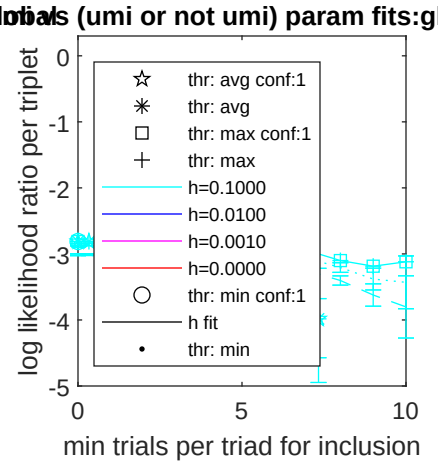

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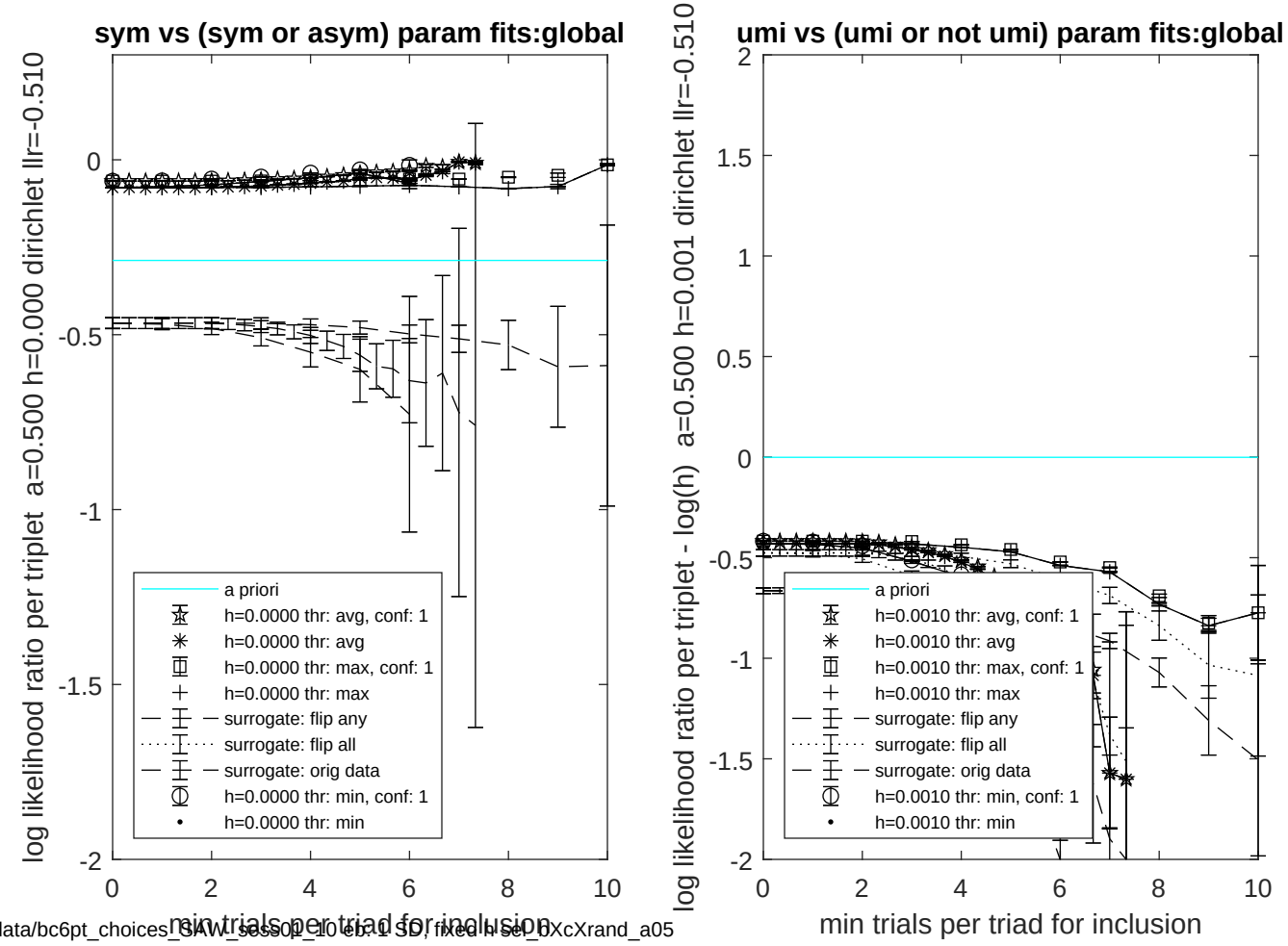

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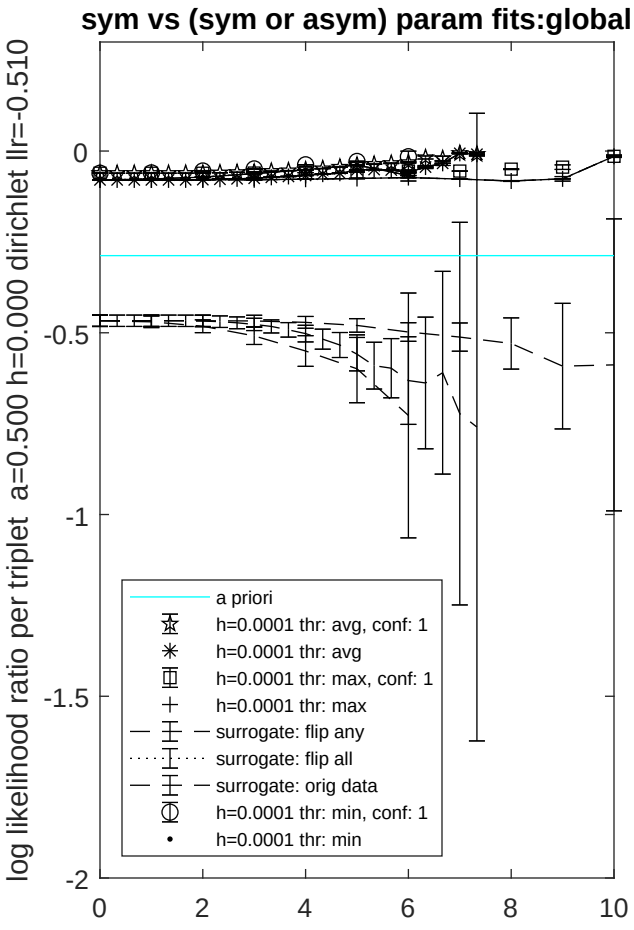

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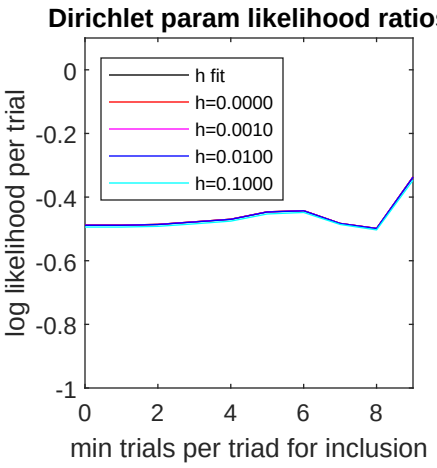

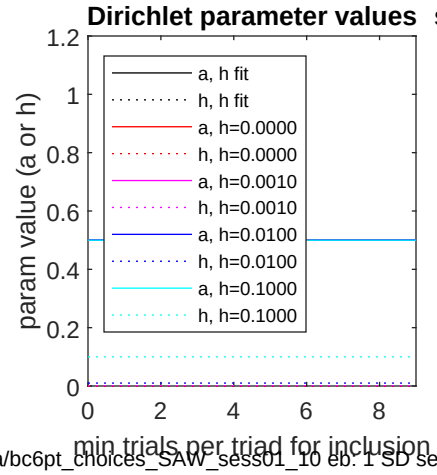

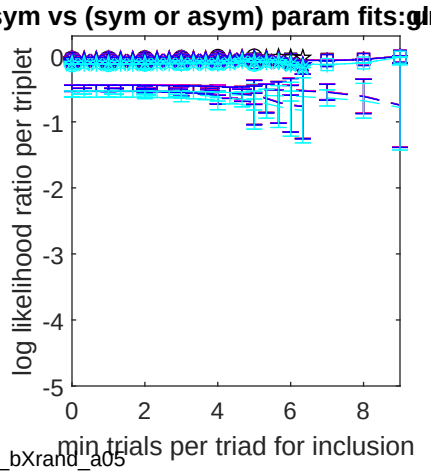

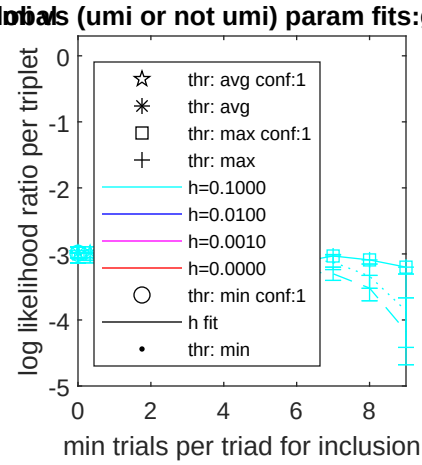

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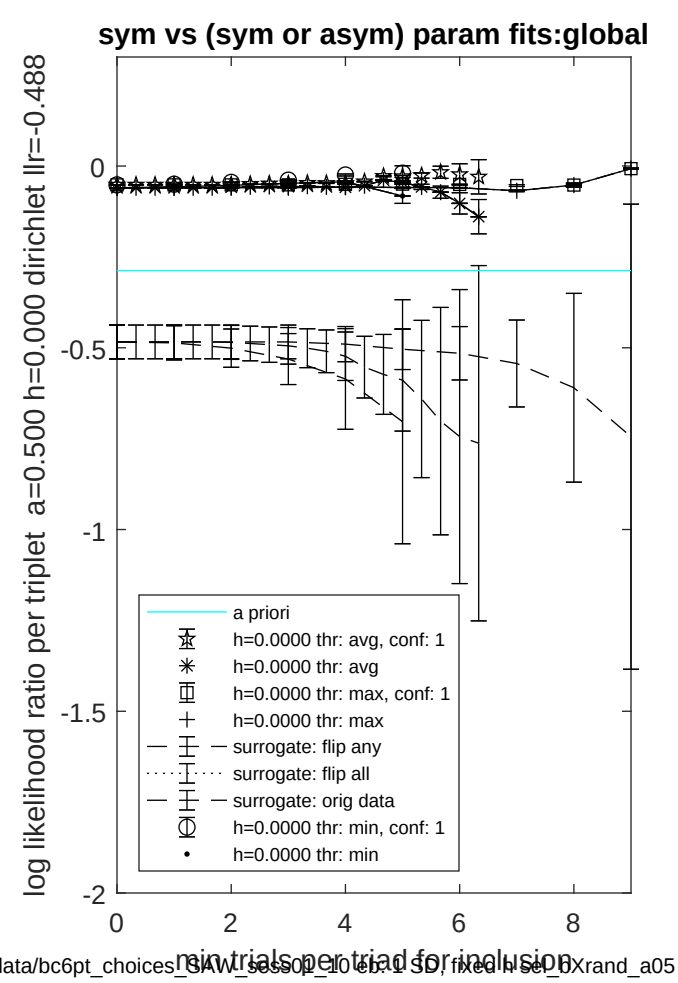

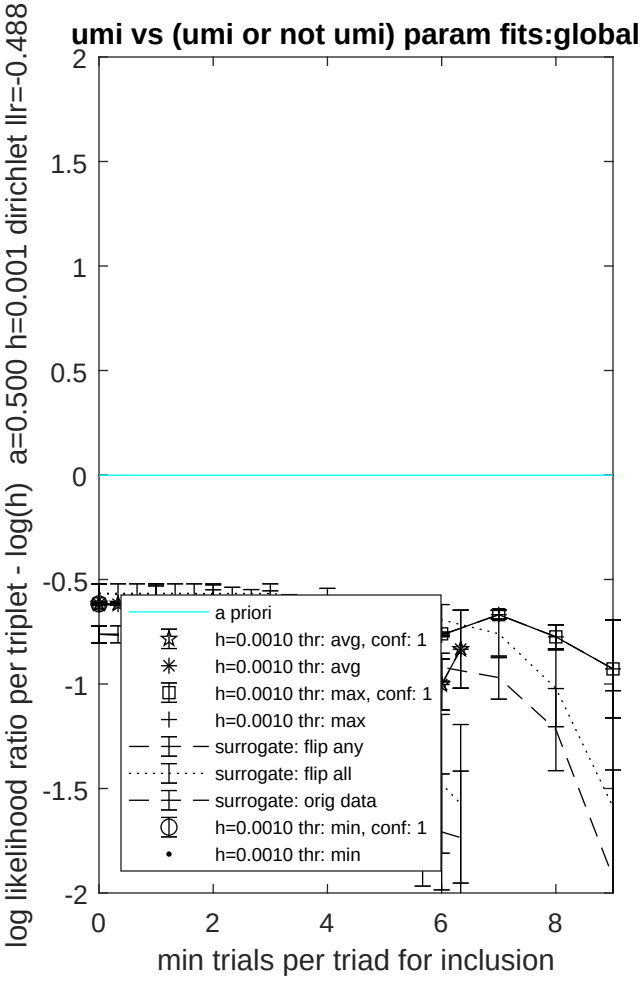

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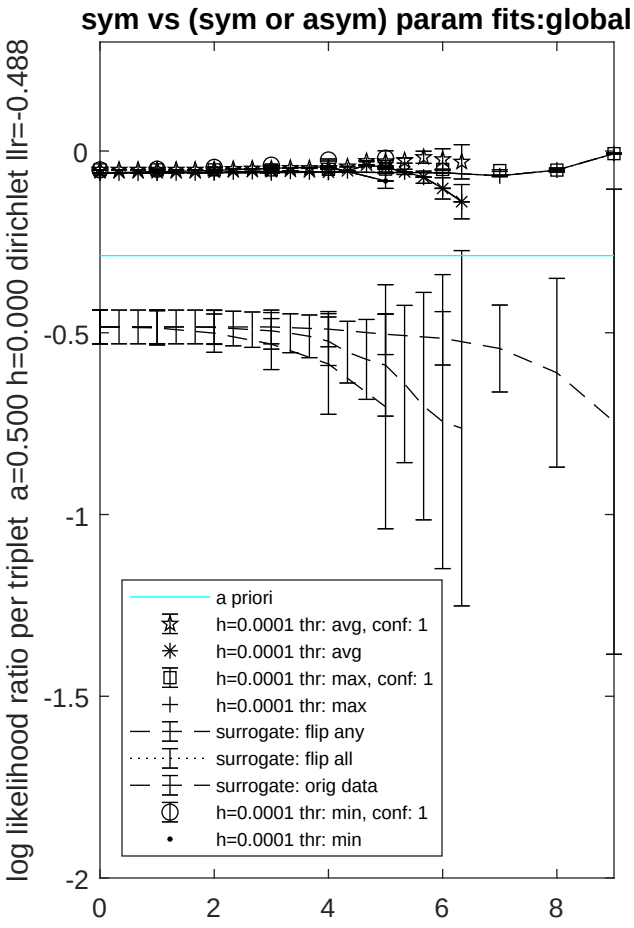

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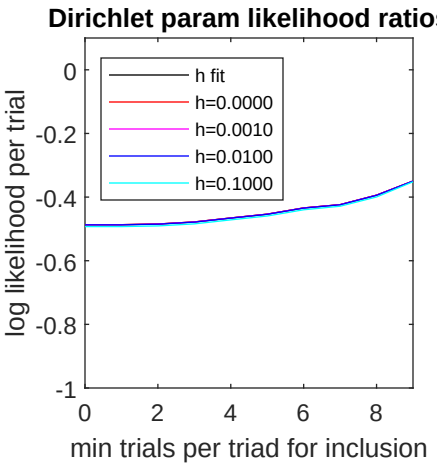

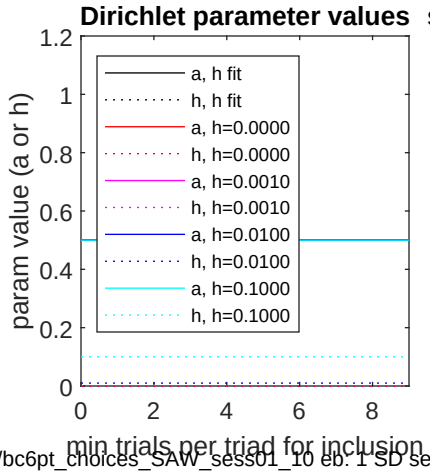

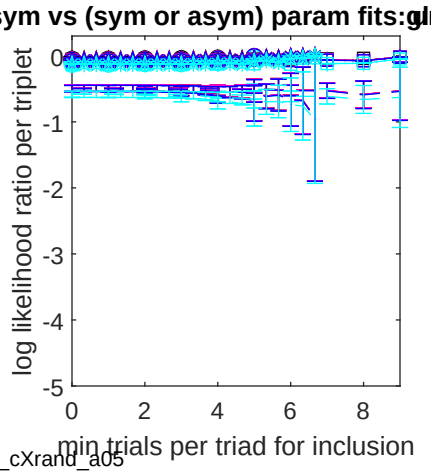

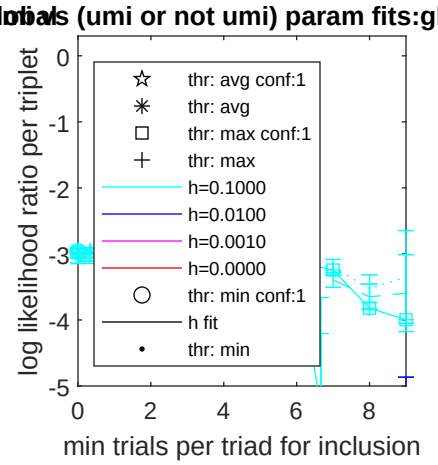

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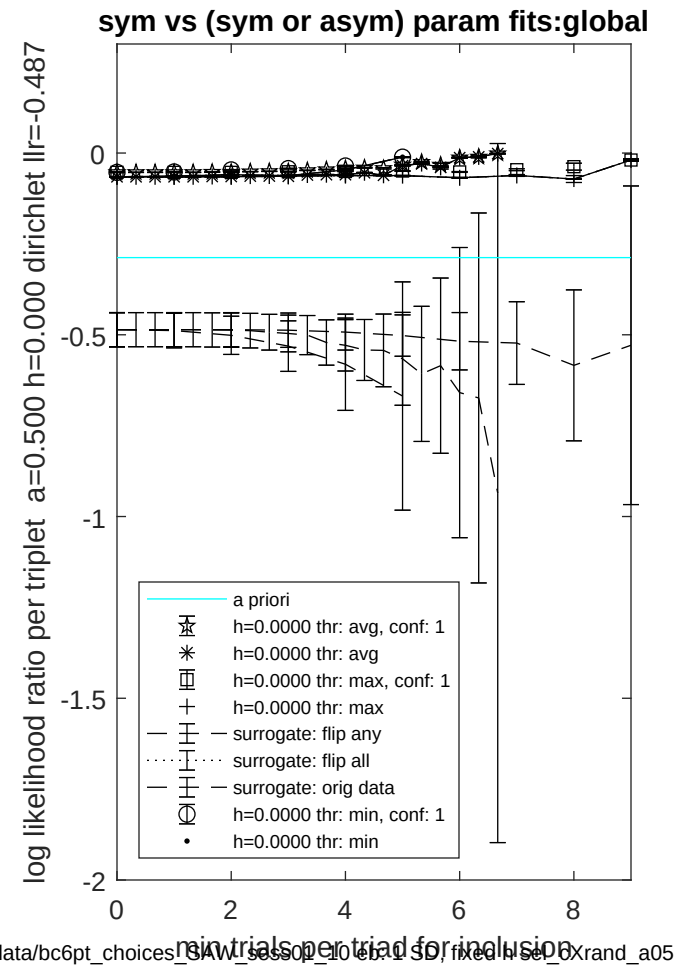

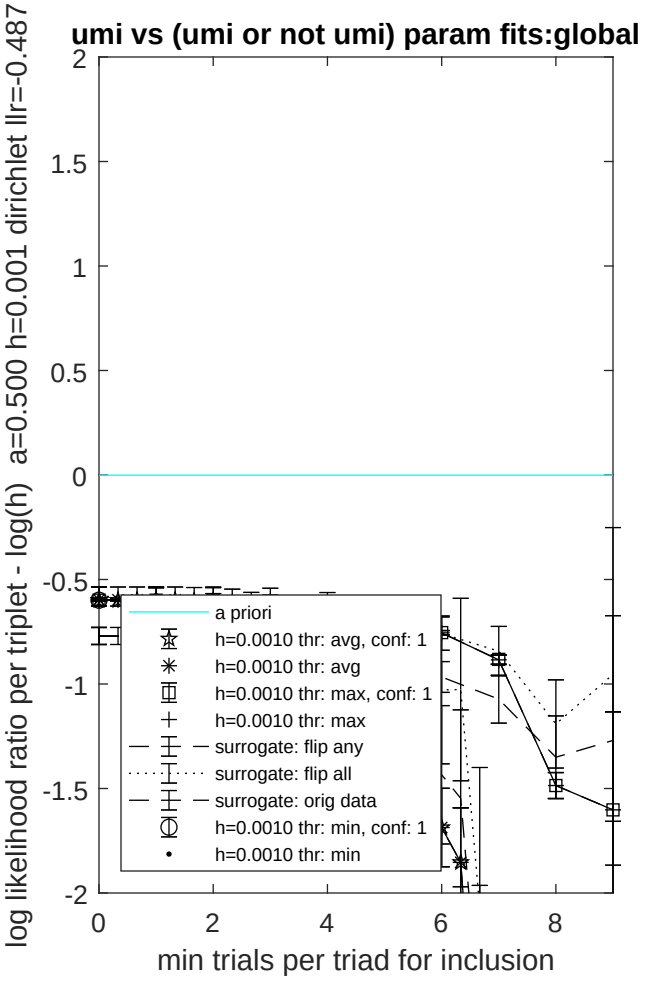

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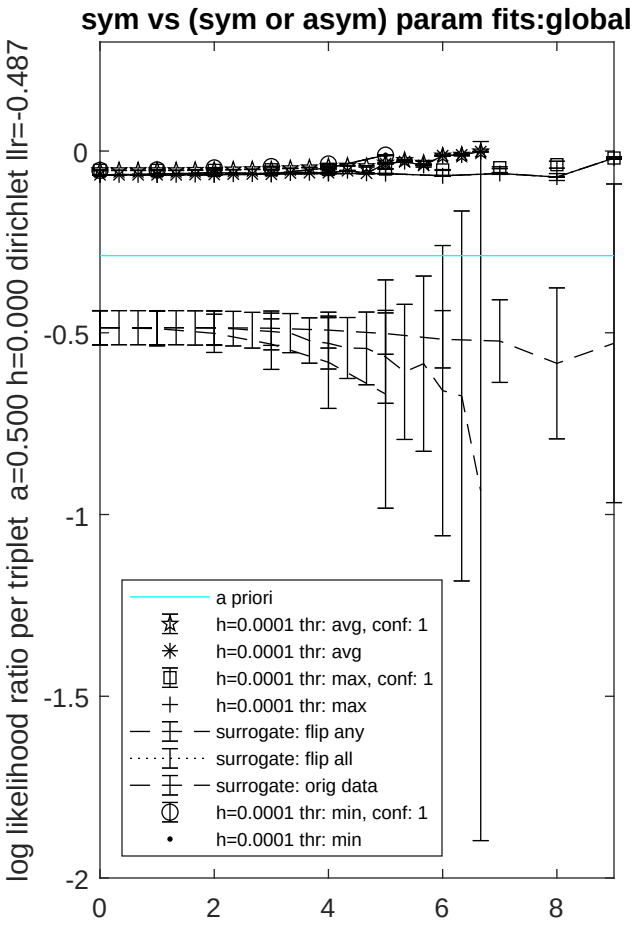
sym vs (sym or asym) param fits:global
log likelihood ratio per triplet a=0.500 h=0.000 dirichlet llr=-0.487
0
-0.5
-1
-1.5
-2
0
2
4
6
8
min trials per triad for inclusion
a priori
h=0.0001 thr: avg, conf: 1
h=0.0001 thr: avg
h=0.0001 thr: max, conf: 1
h=0.0001 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0001 thr: min, conf: 1
h=0.0001 thr: min
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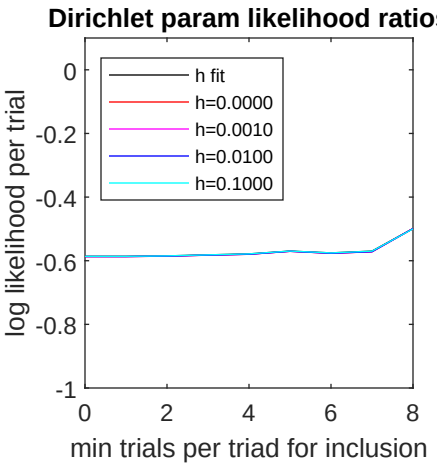

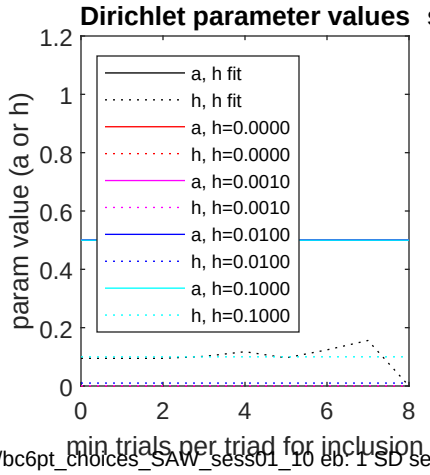

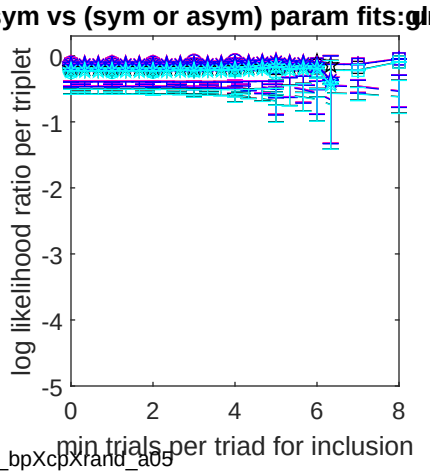

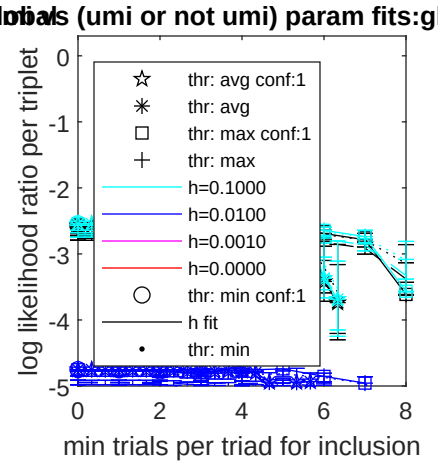

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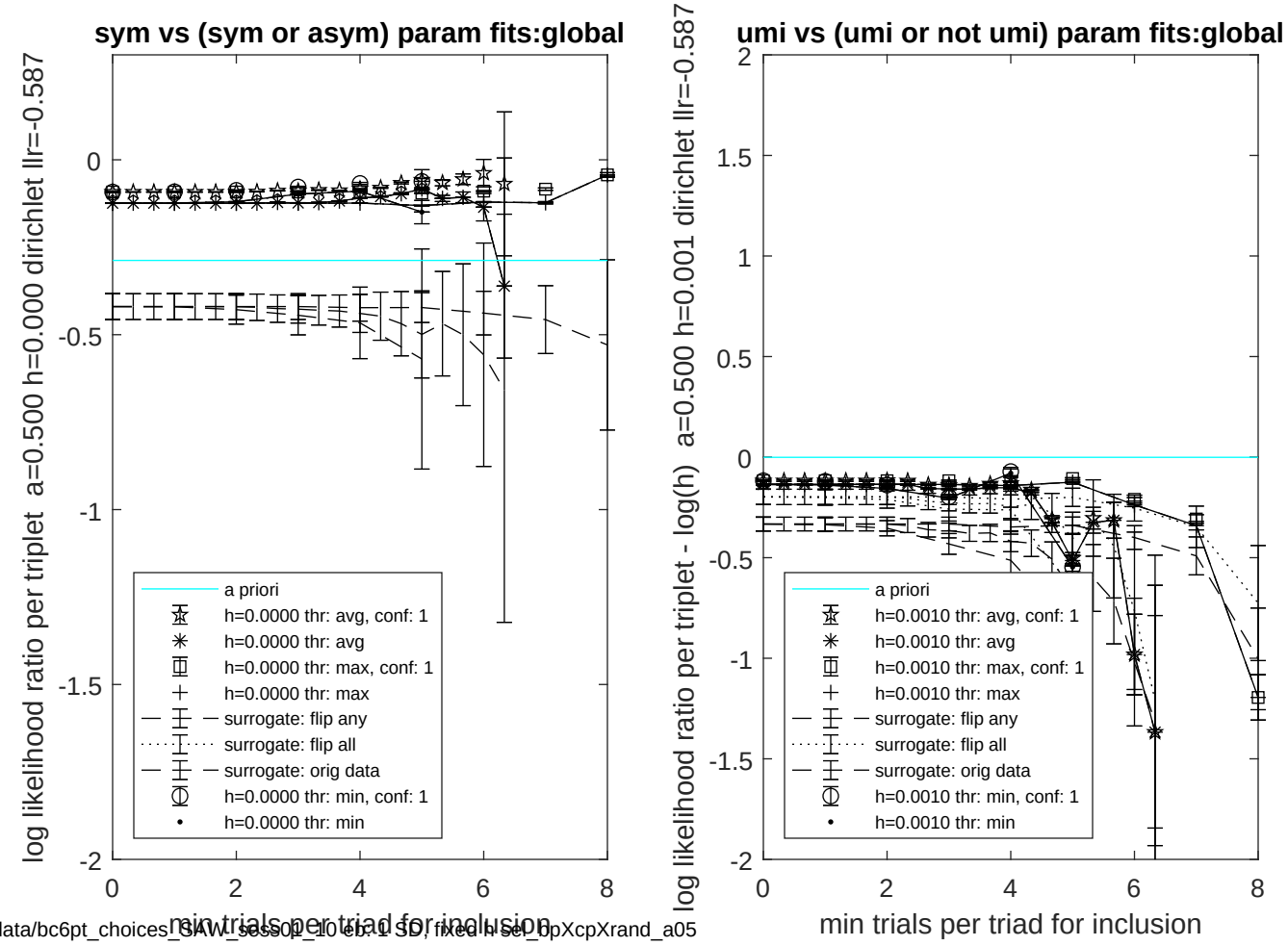

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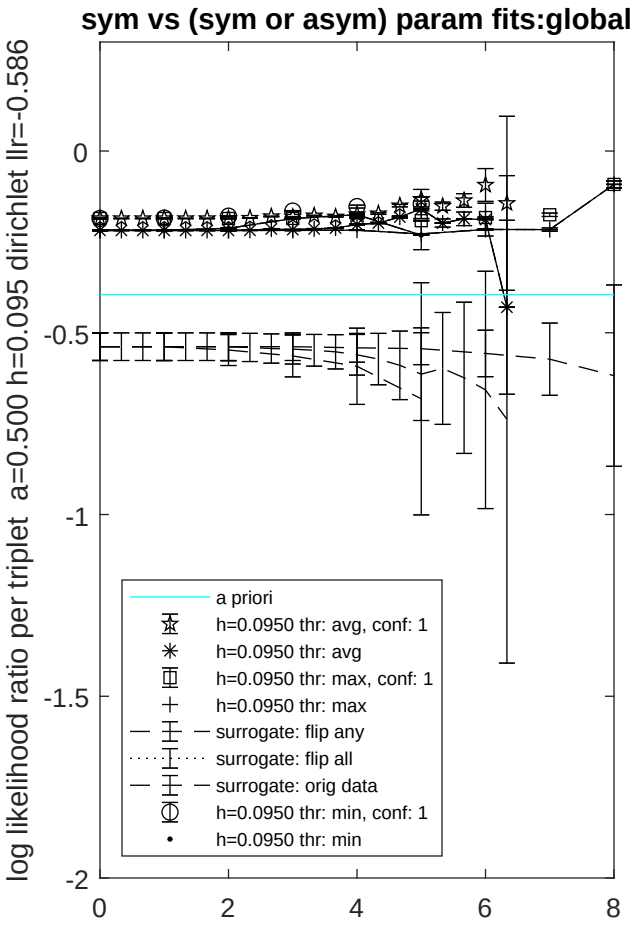

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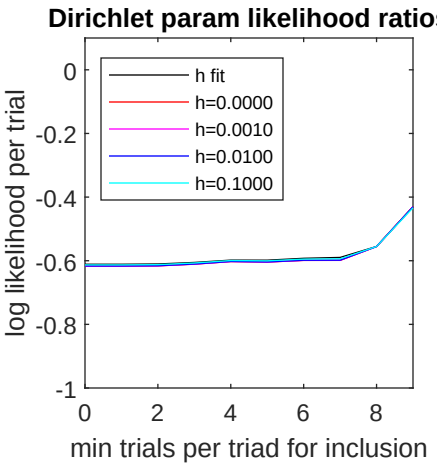

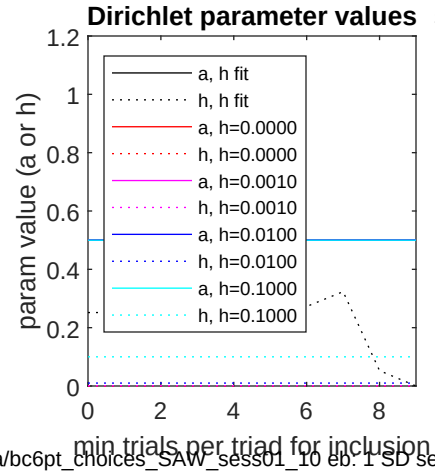

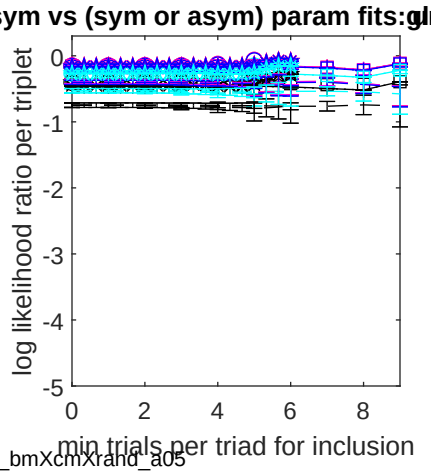

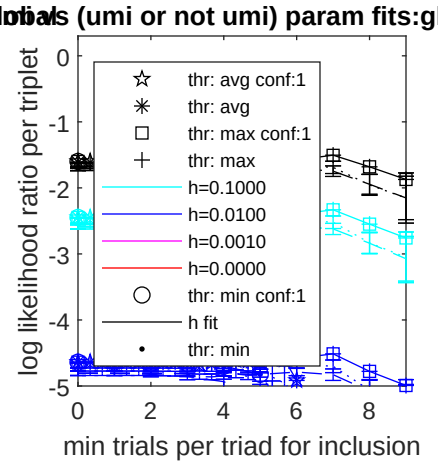

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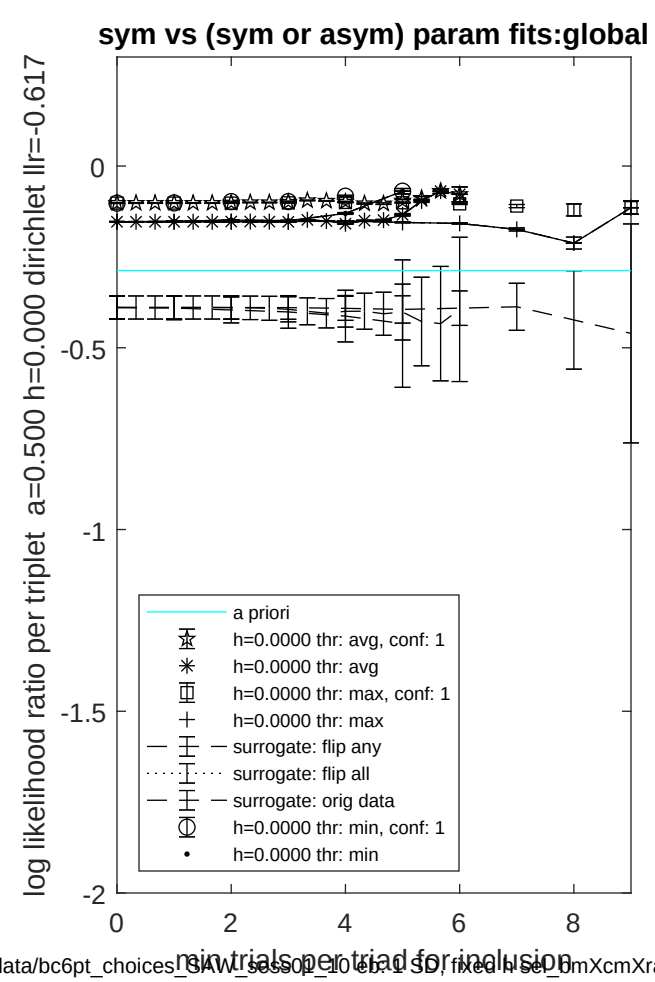

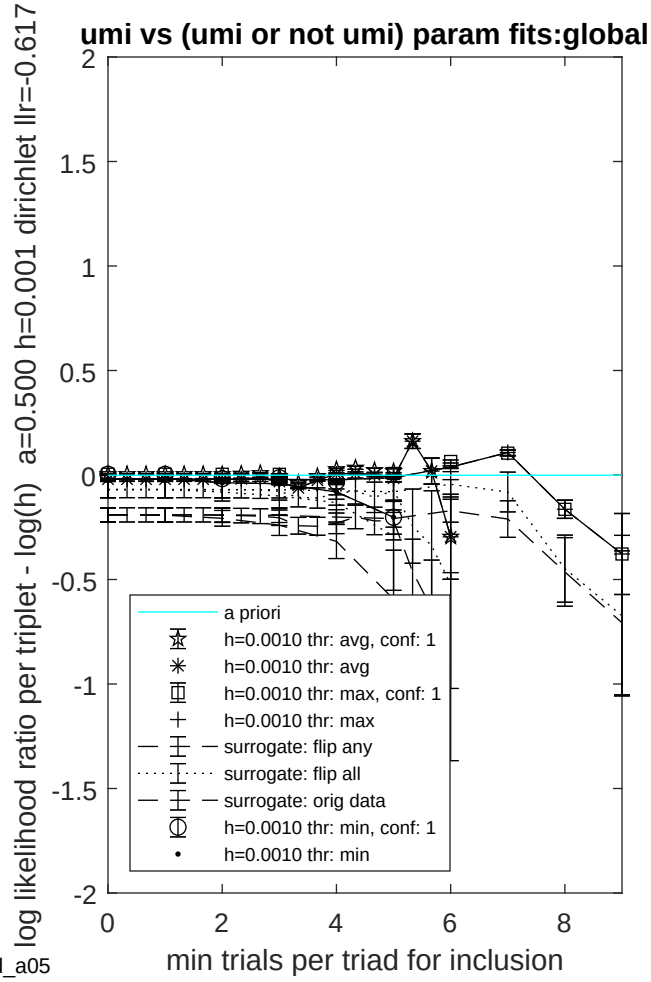

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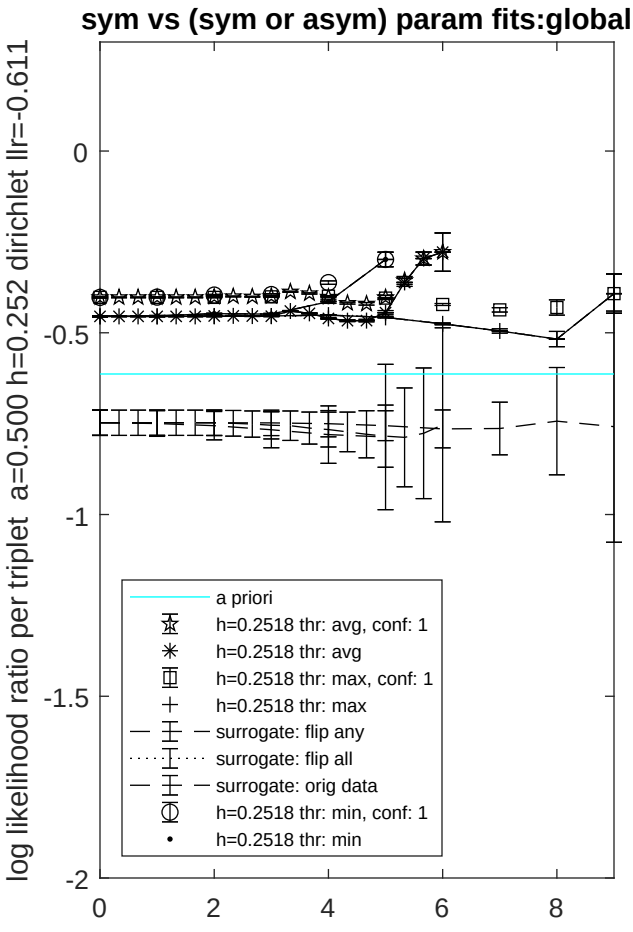

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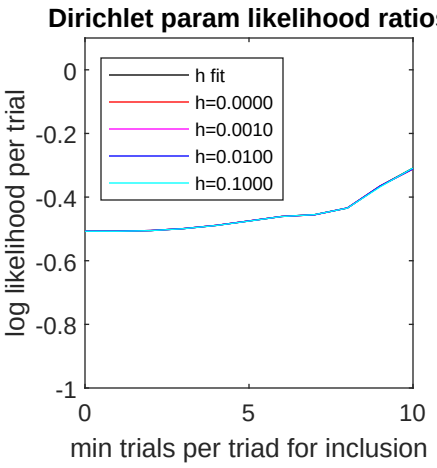

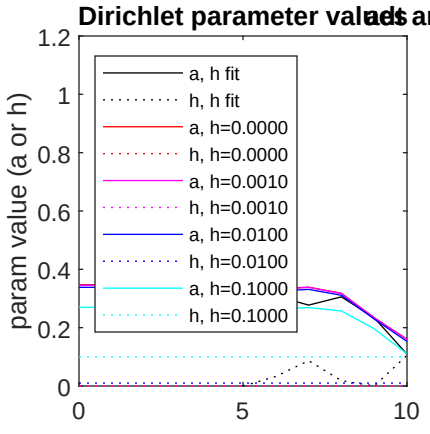

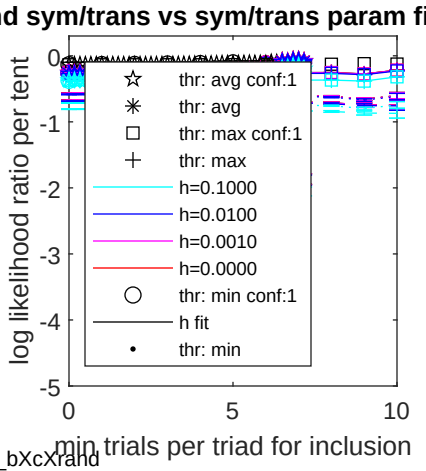

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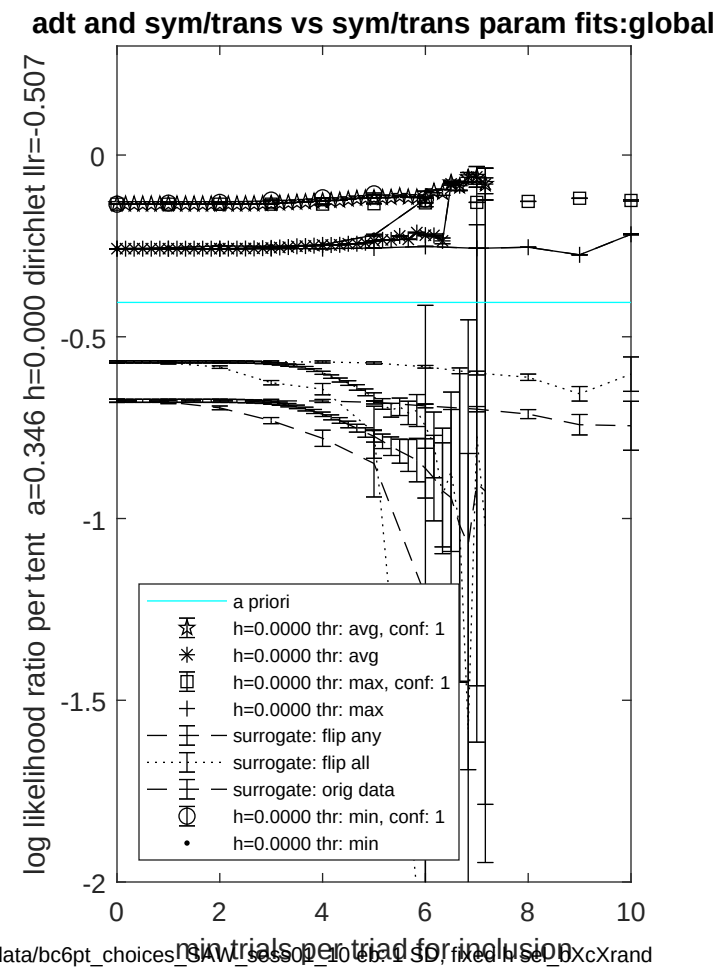
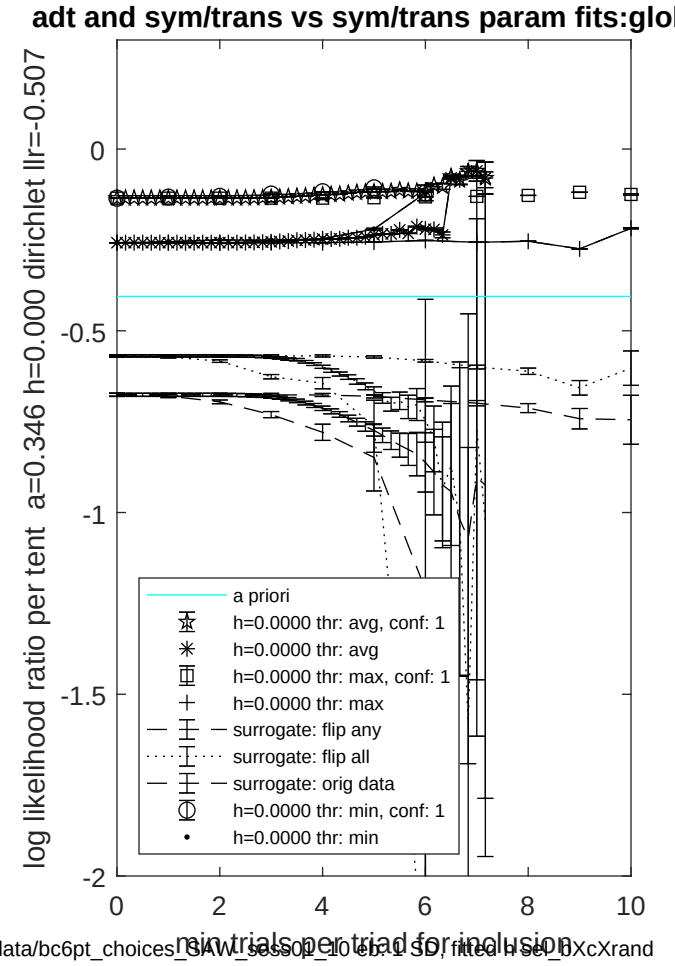
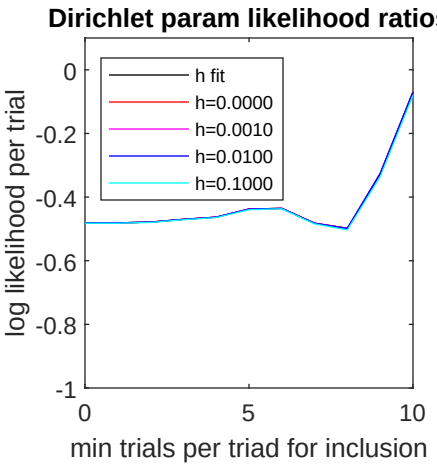
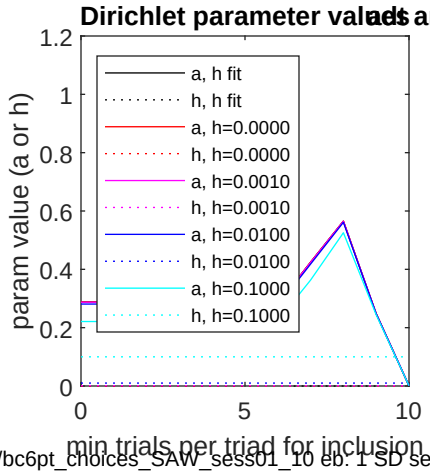
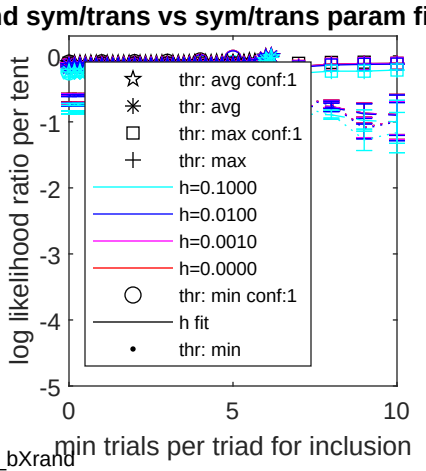

adt and sym/trans vs sym/trans param fits:global
log likelihood ratio per tent a=0.346 h=0.000 dirichlet llr=-0.507
0
-0.5
-1
-1.5
-2
0
2
4
6
8
10
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
./psg_data/bc6pt_choices_SAW_sess0p_10 eb: 1 SD, fixed h sel_bXcXrand

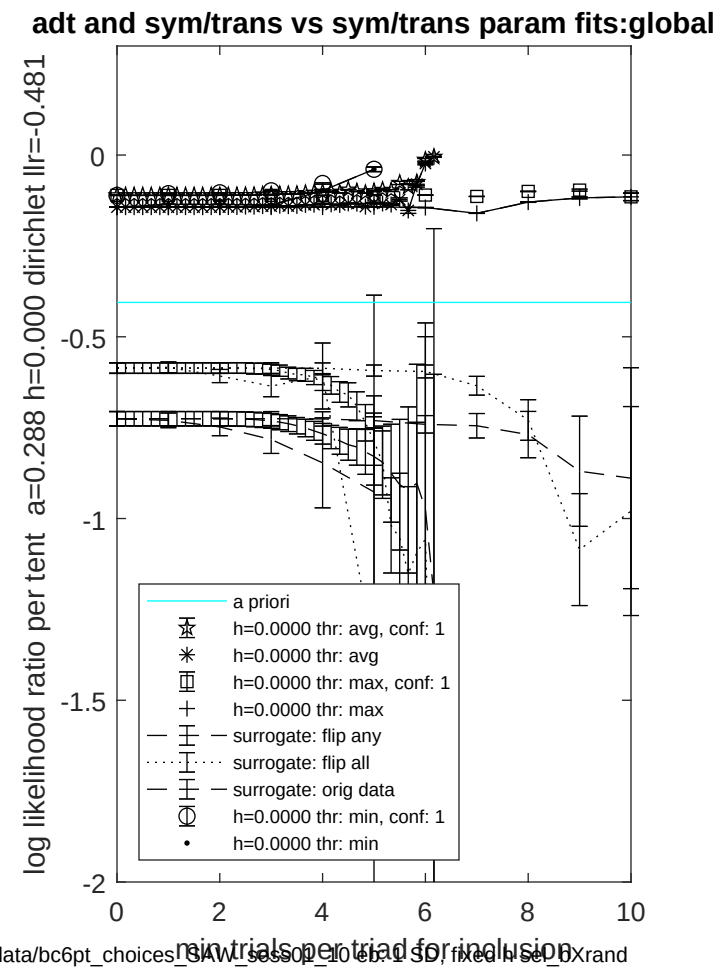

adt and sym/trans vs sym/trans param fits:global
log likelihood ratio per tent a=0.346 h=0.000 dirichlet llr=-0.507
0
-0.5
-1
-1.5
-2
0
2
4
6
8
10
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
./psg_data/bc6pt_choices_SAW_sess0p_10 eb: 1 SD, fitted h sel_bXcXrand

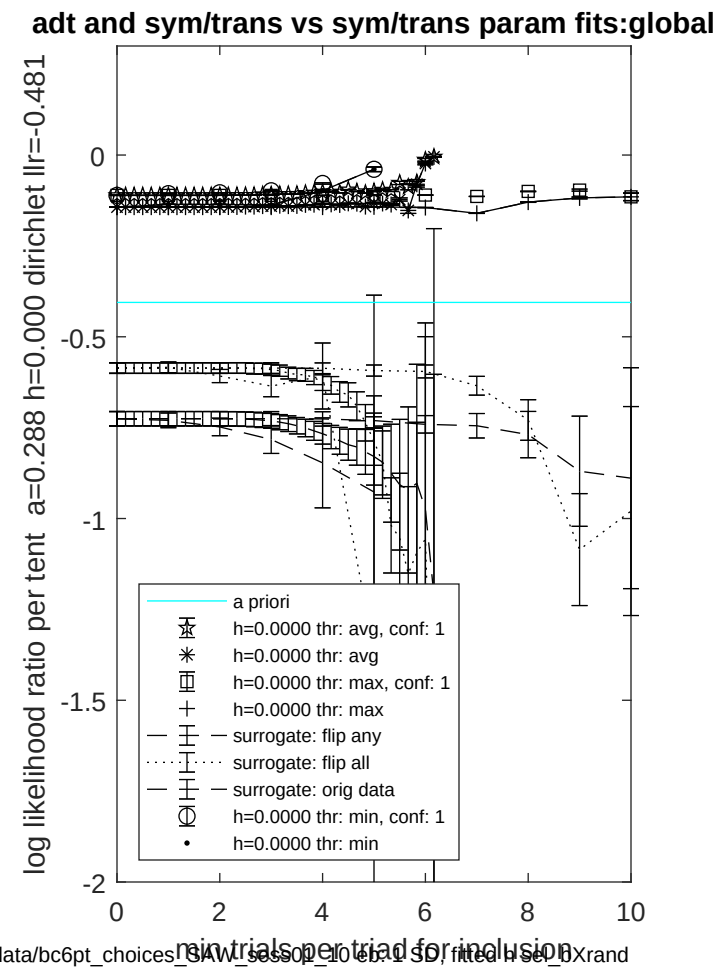


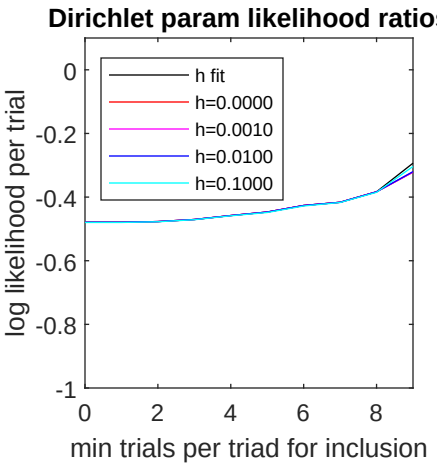


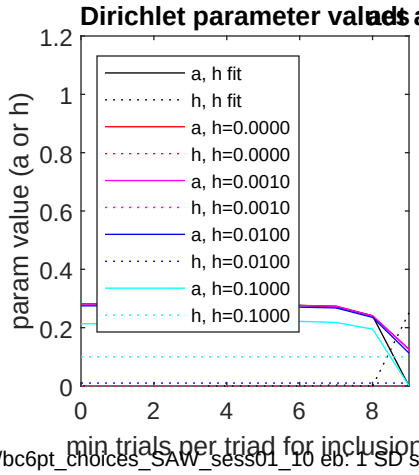


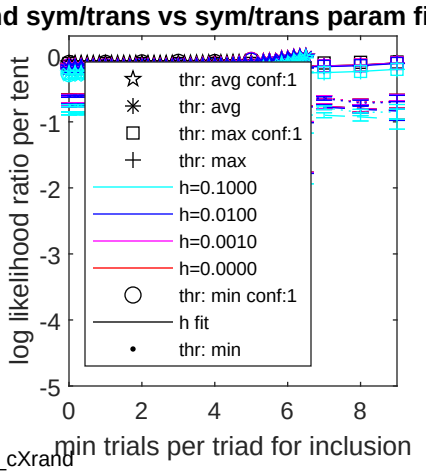
./psg_data/bc6pt_choices_SAW_sess01_10 eb: 1 SD sel_bXrand


adt and sym/trans vs sym/trans param fits:global
log likelihood ratio per tent a=0.288 h=0.000 dirichlet llr=-0.481
0
-0.5
-1
-1.5
-2
0
2
4
6
8
10
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
./psg_data/bc6pt_choices_SAW_sess01_10 eb: 1 SD, fixed h sel_bXrand

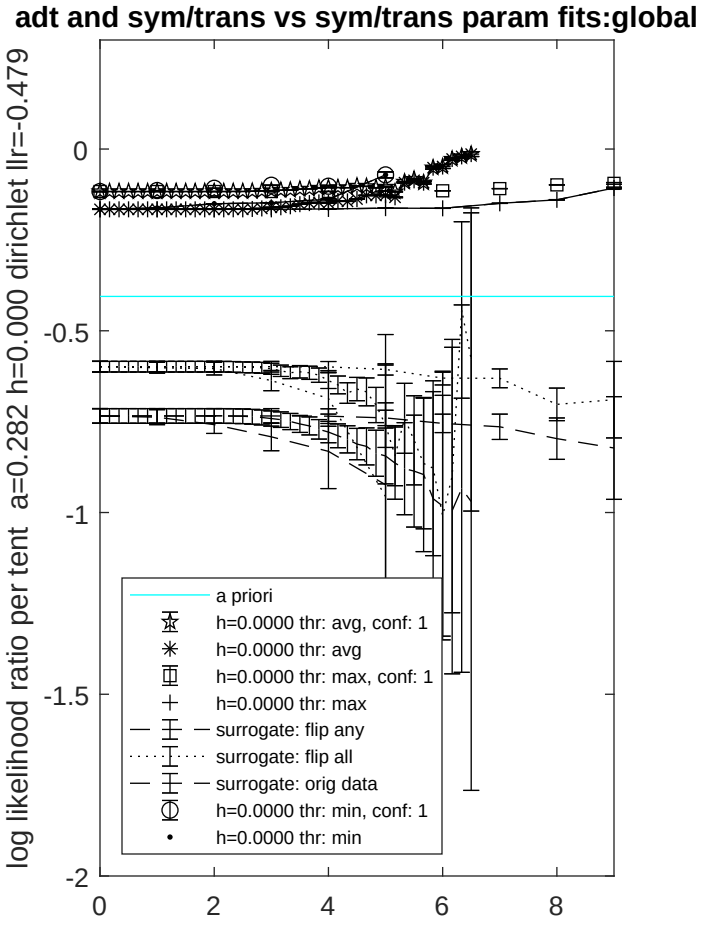

adt and sym/trans vs sym/trans param fits:global
log likelihood ratio per tent a=0.288 h=0.000 dirichlet llr=-0.481
0
-0.5
-1
-1.5
-2
0
2
4
6
8
10
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
./psg_data/bc6pt_choices_SAW_sess01_10 eb: 1 SD, fitted h sel_bXrand

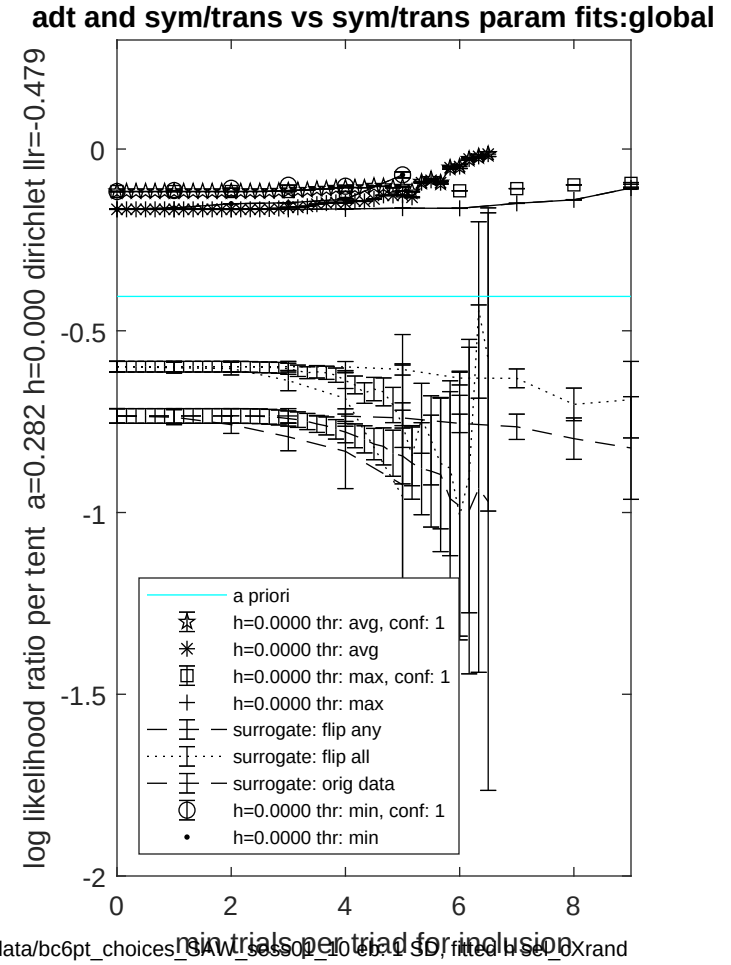


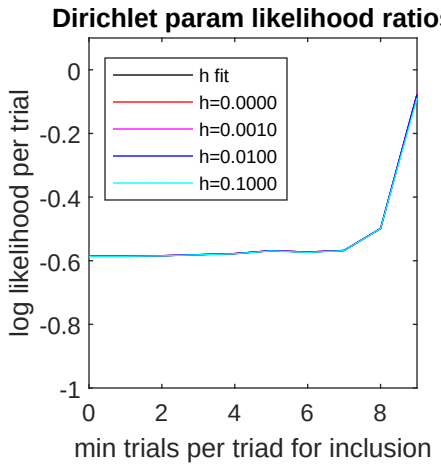


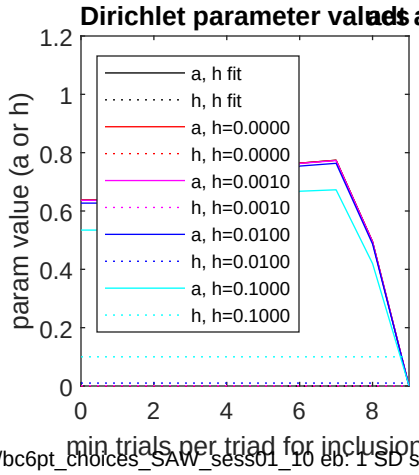


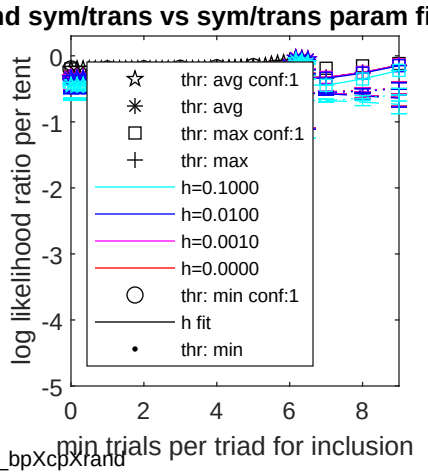
./psg_data/bc6pt_choices_SAW_sess01_10 eb: 1 SD sel_cXrand


adt and sym/trans vs sym/trans param fits:global
log likelihood ratio per tent a=0.282 h=0.000 dirichlet llr=-0.479
0
-0.5
-1
-1.5
-2
0
2
4
6
8
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
./psg_data/bc6pt_choices_SAW_sess01_10 eb: 1 SD, fixed h sel_0Xrand


adt and sym/trans vs sym/trans param fits:global
log likelihood ratio per tent a=0.282 h=0.000 dirichlet llr=-0.479
0
-0.5
-1
-1.5
-2
0
2
4
6
8
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
./psg_data/bc6pt_choices_SAW_sess01_10 eb: 1 SD, fitted h sel_0Xrand







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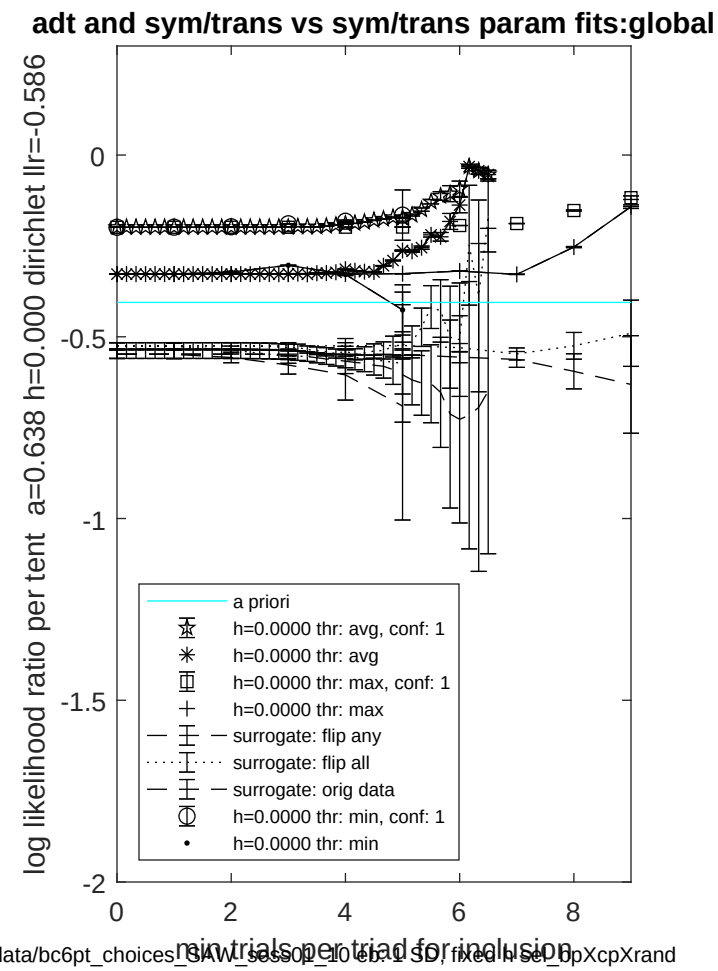
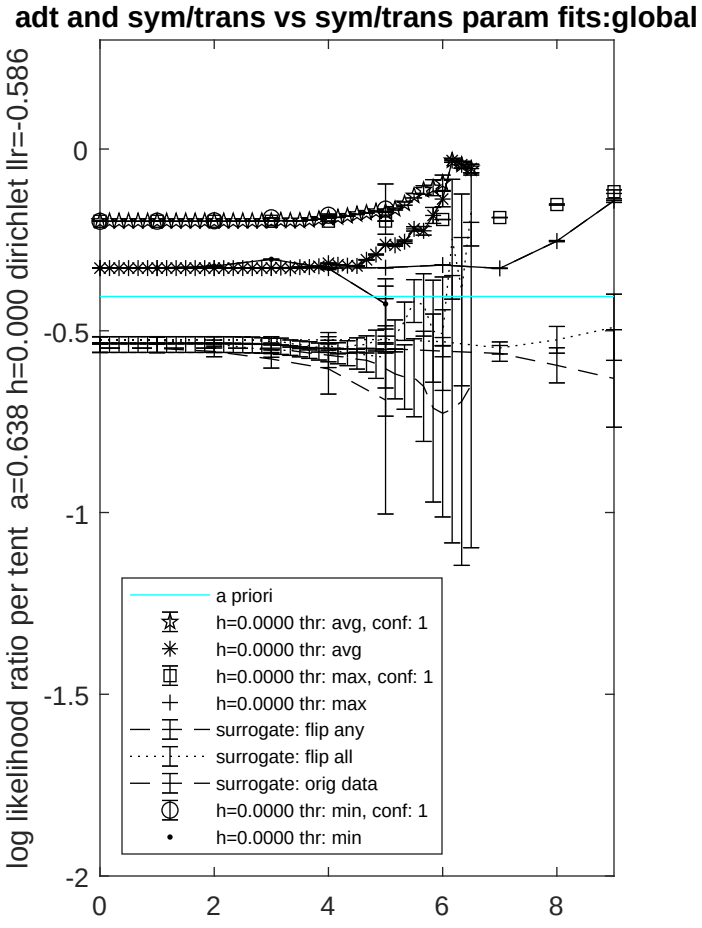
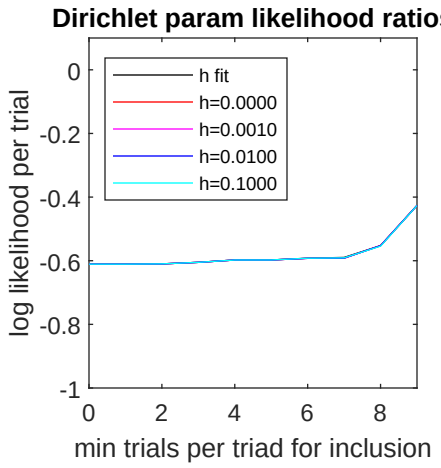
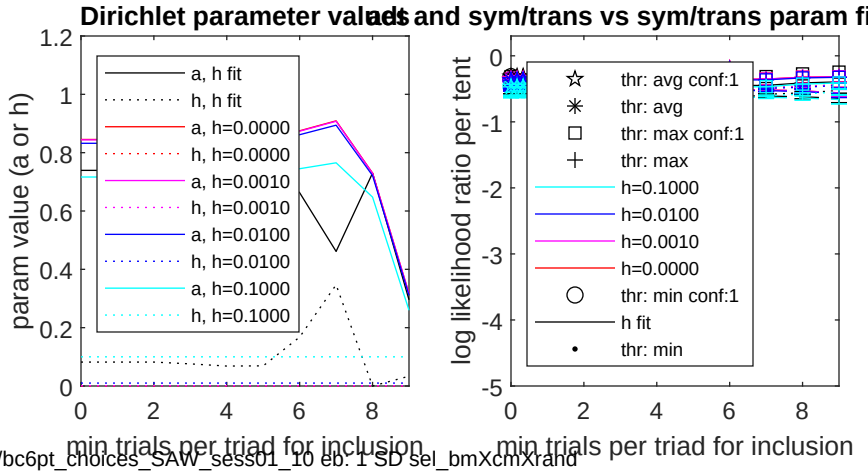

adt and sym/trans vs sym/trans param fits:global
log likelihood ratio per tent a=0.638 h=0.000 dirichlet llr=-0.586
0
-0.5
-1
-1.5
-2
0
2
4
6
8
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
./psg_data/bc6pt_choices_SAW_sess01_10 eb: 1 SD, fixed h sel_bpXcpXrand


adt and sym/trans vs sym/trans param fits:global
log likelihood ratio per tent a=0.638 h=0.000 dirichlet llr=-0.586
0
-0.5
-1
-1.5
-2
0
2
4
6
8
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
./psg_data/bc6pt_choices_SAW_sess01_10 eb: 1 SD, fitted h sel_bpXcpXrand





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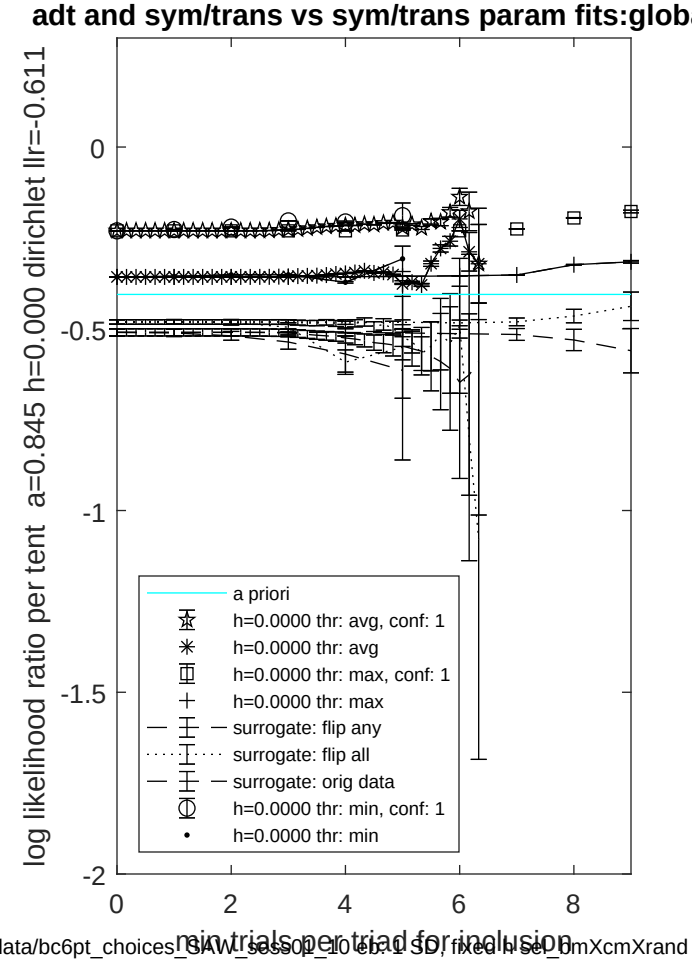

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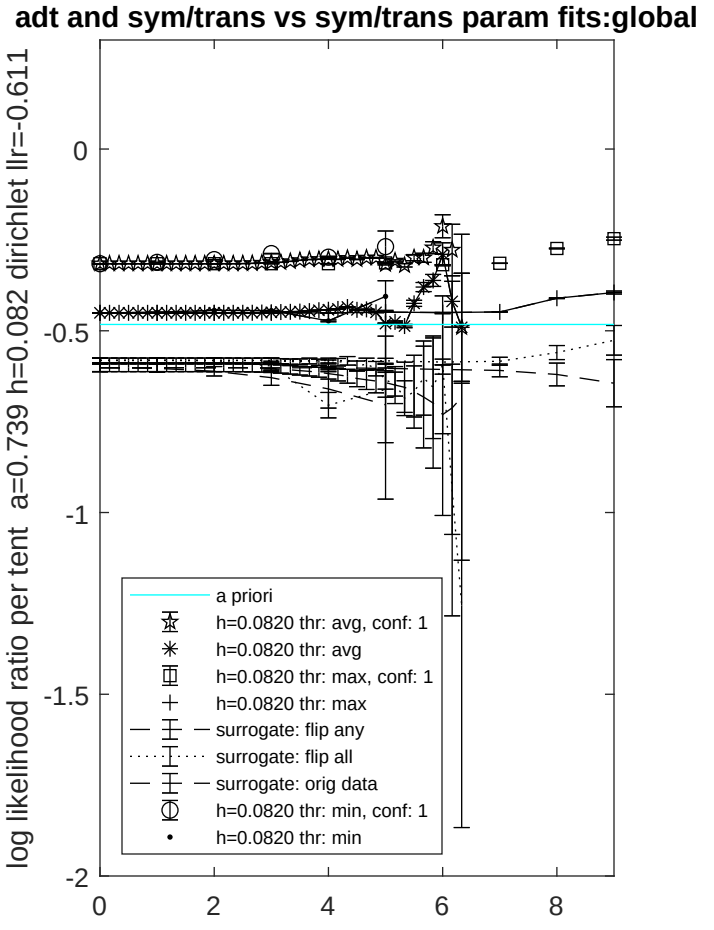

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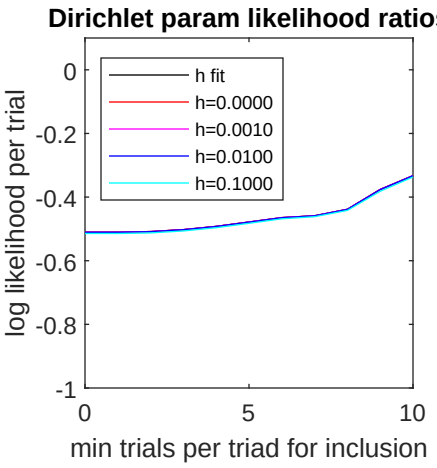

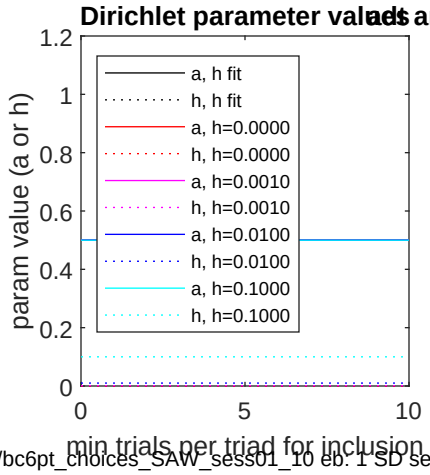

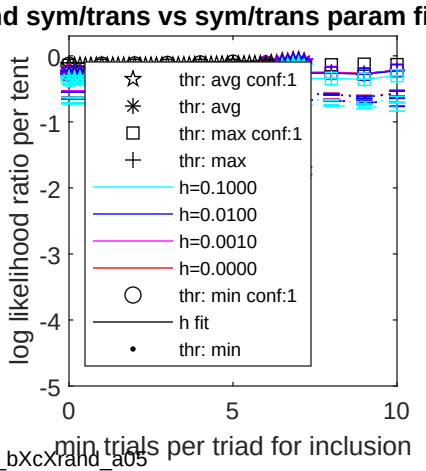

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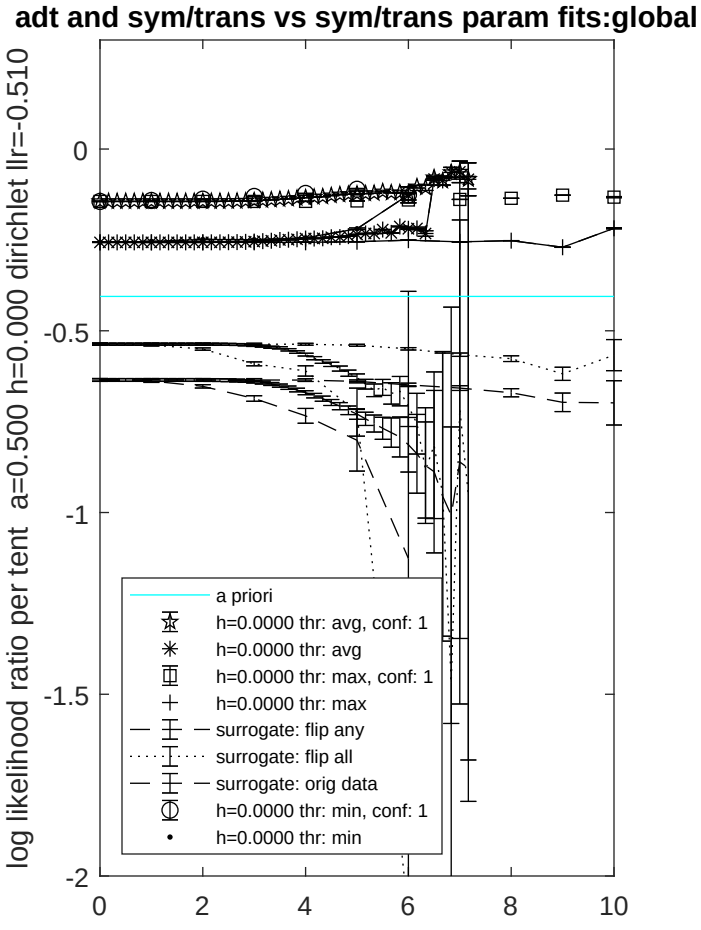

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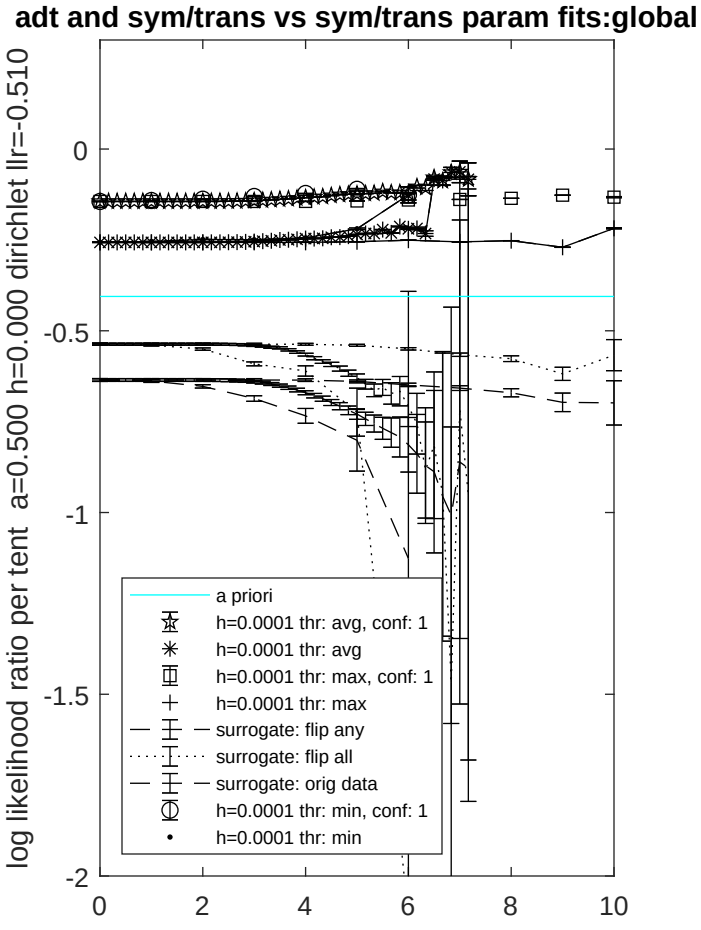

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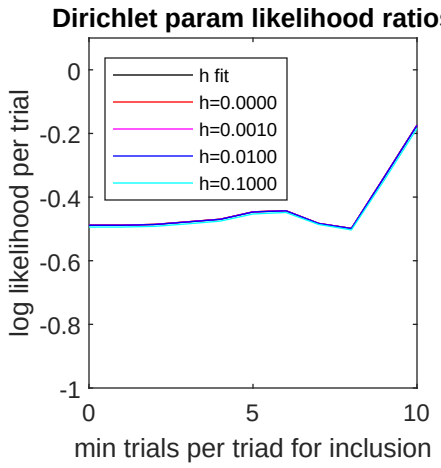

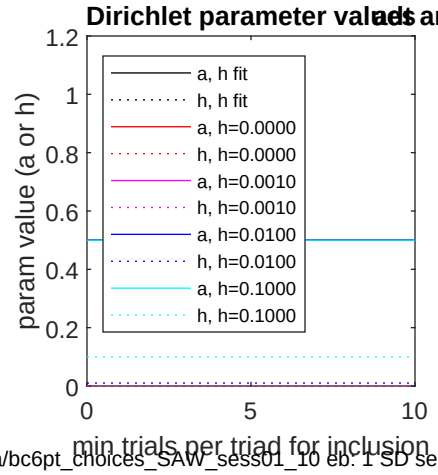

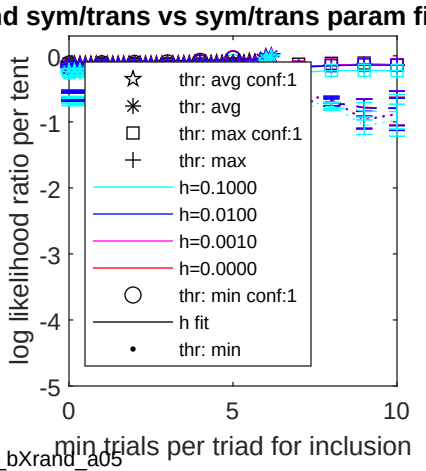

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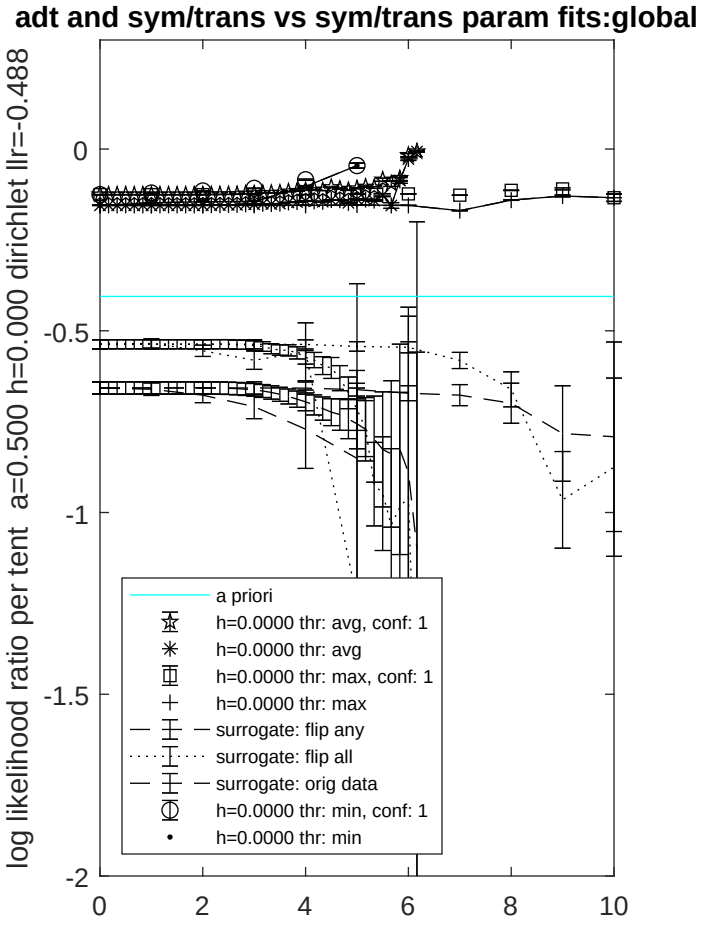

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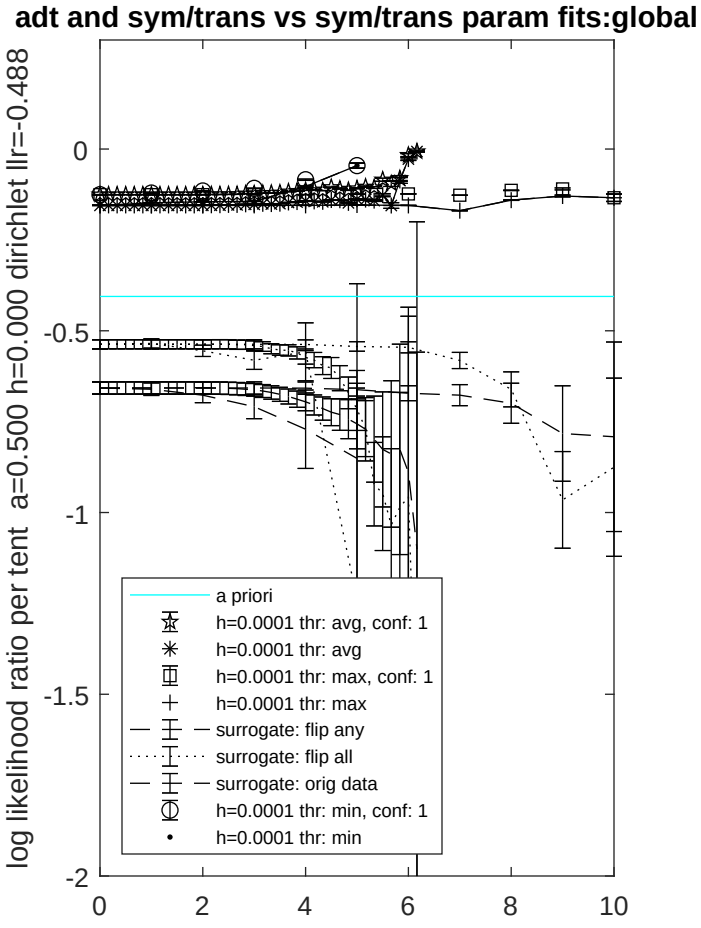

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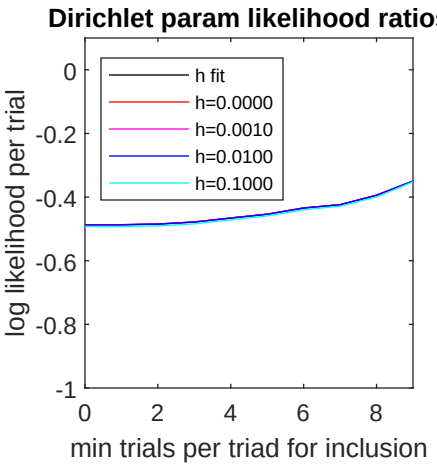

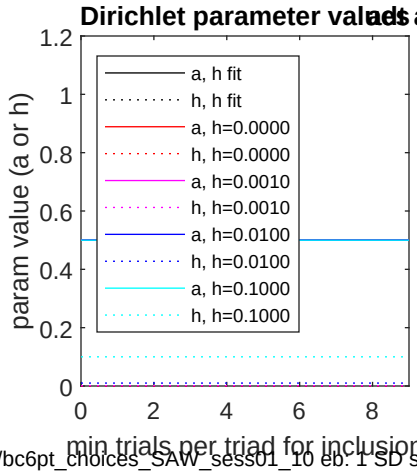

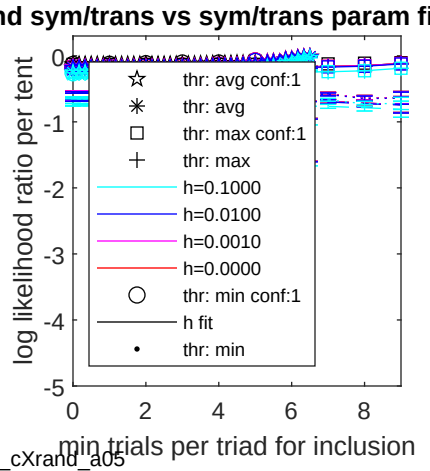

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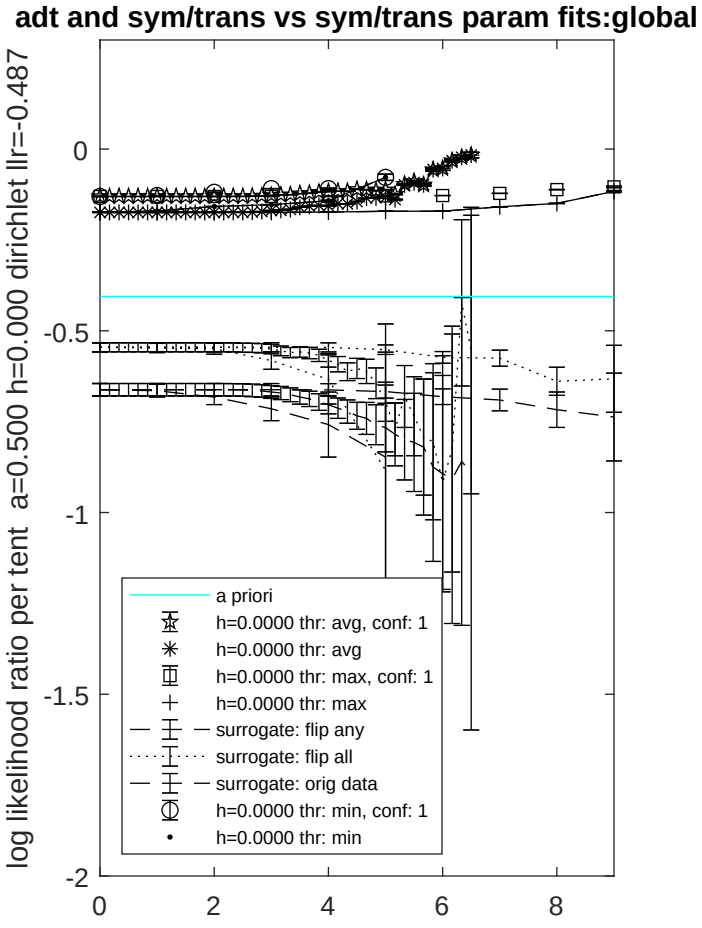

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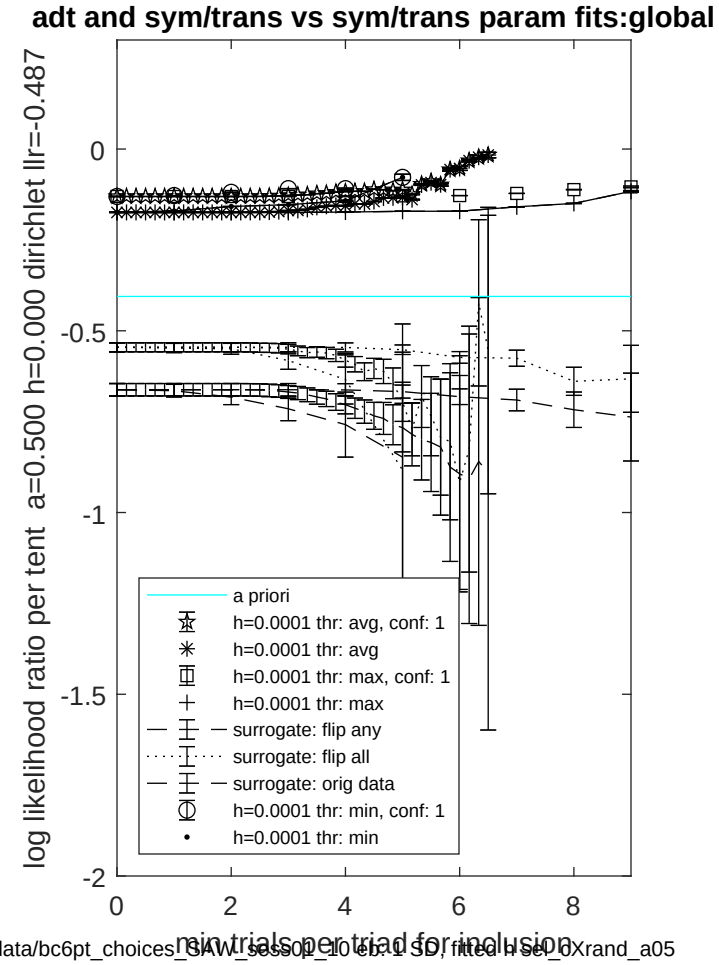
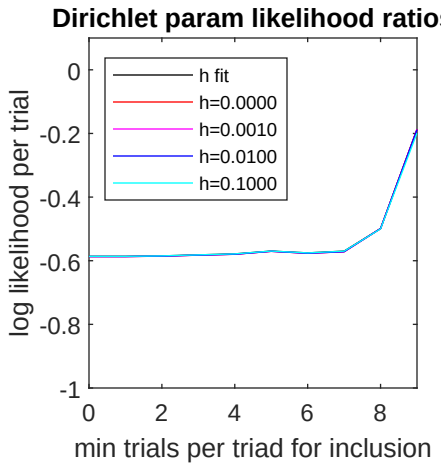
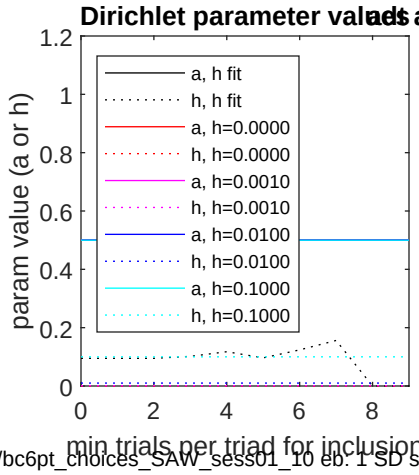
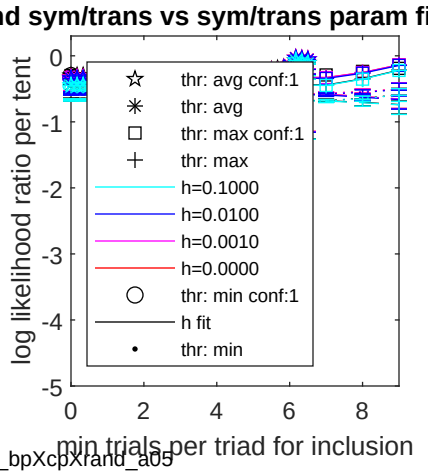


./psg_data/bc6pt_choices_SAW_sess01_10 eb:1 SD, fitted h set_0Xrand_a05







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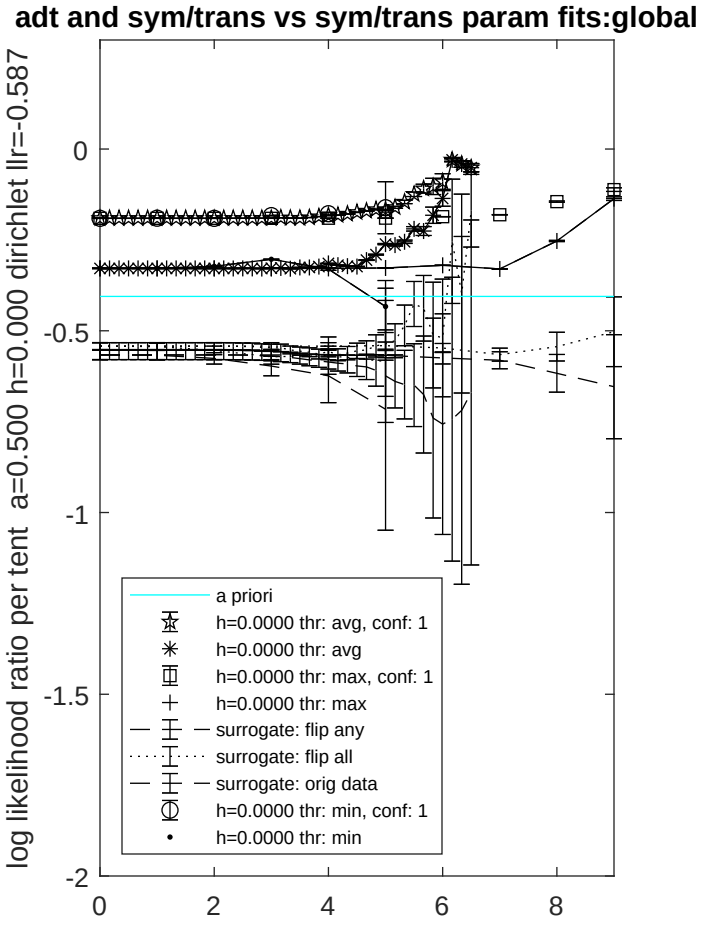
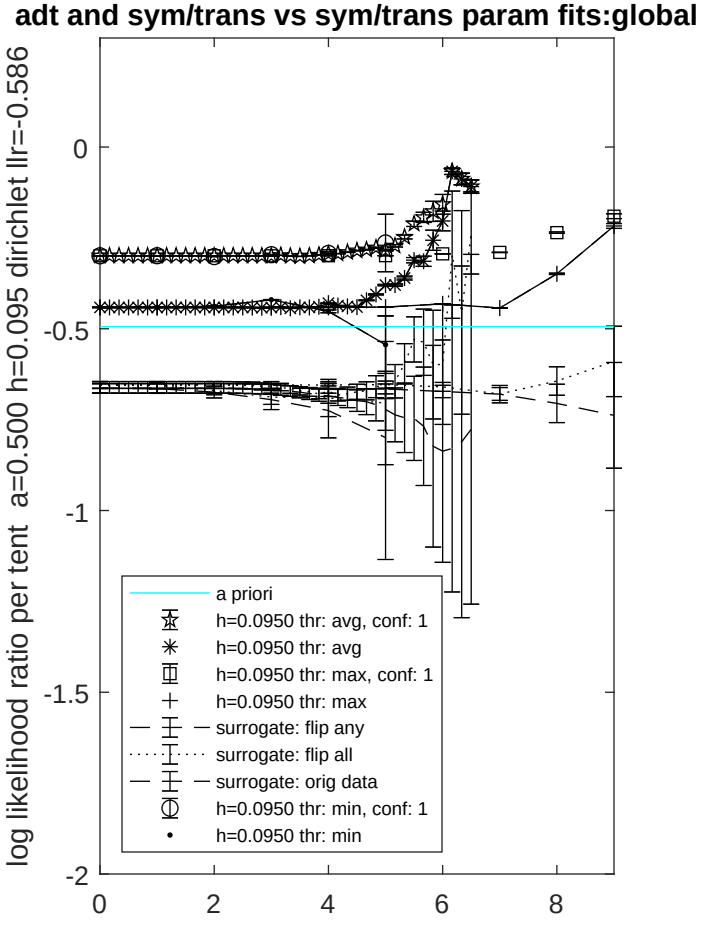
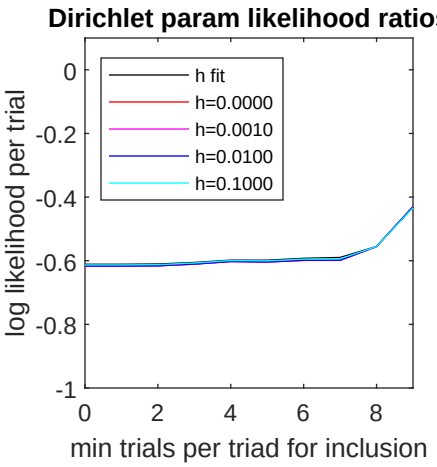
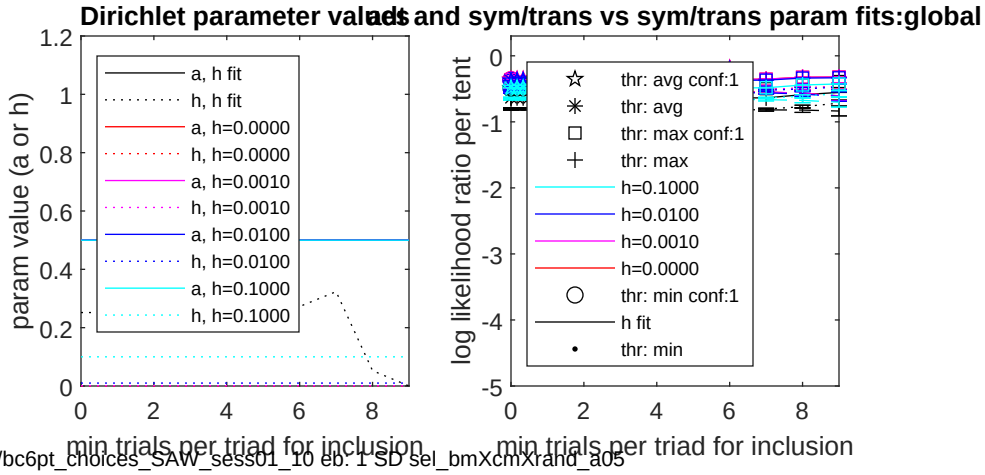


./psg_data/bc6pt_choices_SAW_sess01_10 eb:1 SD, fixed h set_6pXcpXrand_a05


adt and sym/trans vs sym/trans param fits:global
log likelihood ratio per tent a=0.500 h=0.095 dirichlet llr=-0.586
0
-0.5
-1
-1.5
-2
0
2
4
6
8
min trials per triad for inclusion
a priori
h=0.0950 thr: avg, conf: 1
h=0.0950 thr: avg
h=0.0950 thr: max, conf: 1
h=0.0950 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0950 thr: min, conf: 1
h=0.0950 thr: min
./psg_data/bc6pt_choices_SAW_sess01_10 eb: 1 SD, fitted h sel_bpXcpXrand_a05





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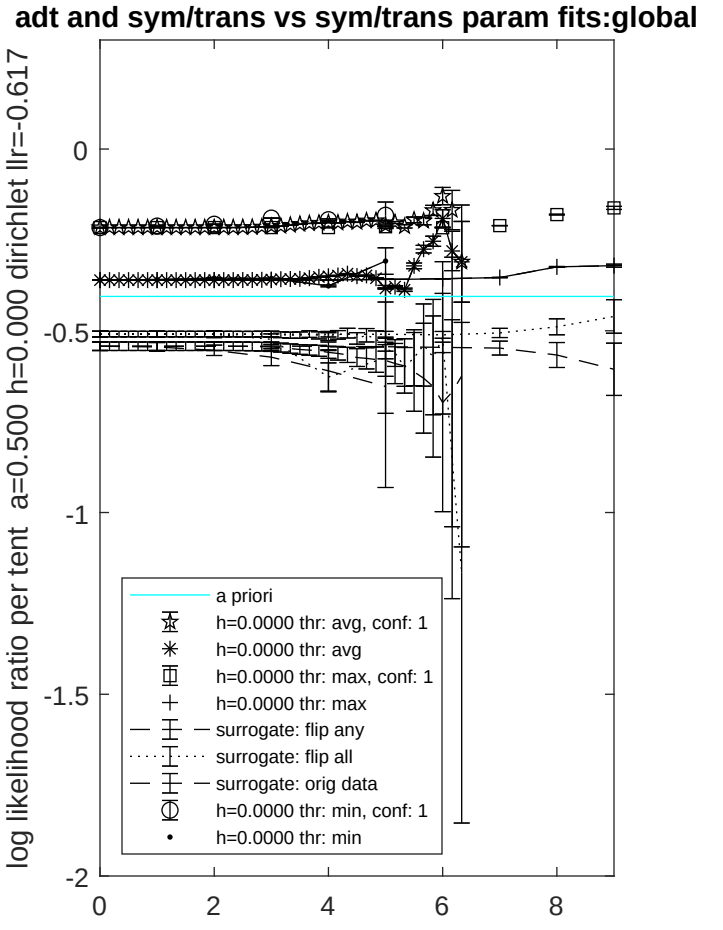

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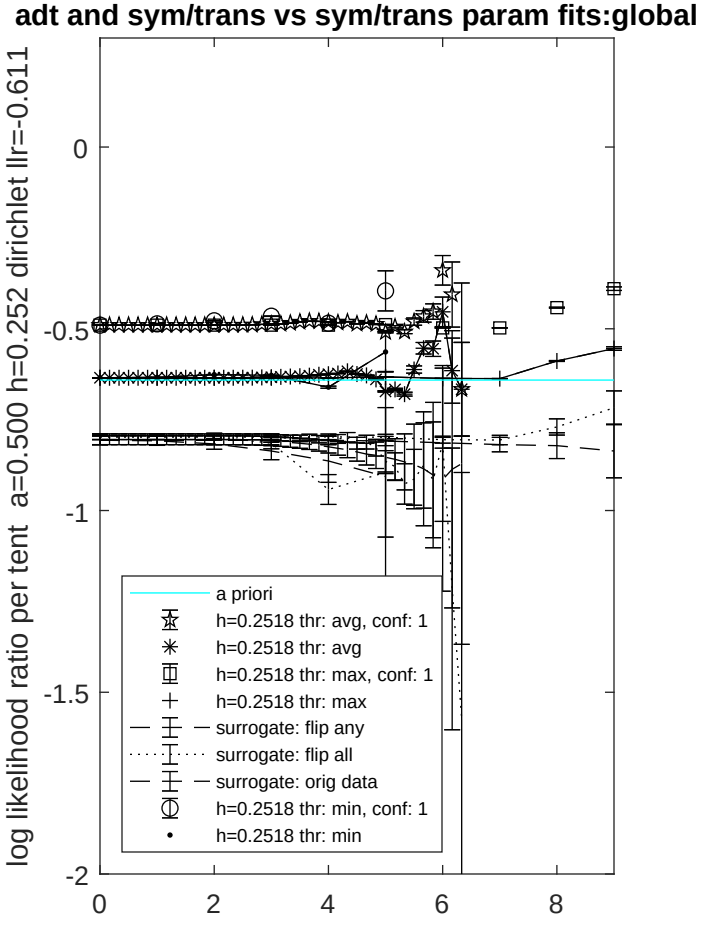

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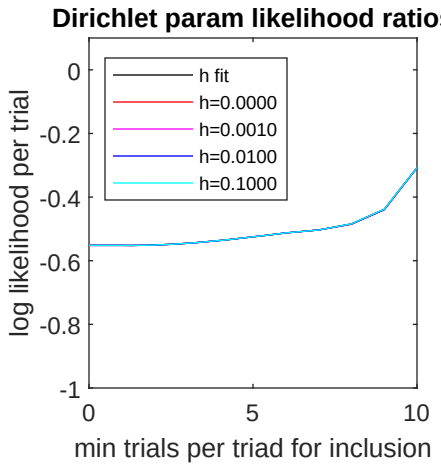

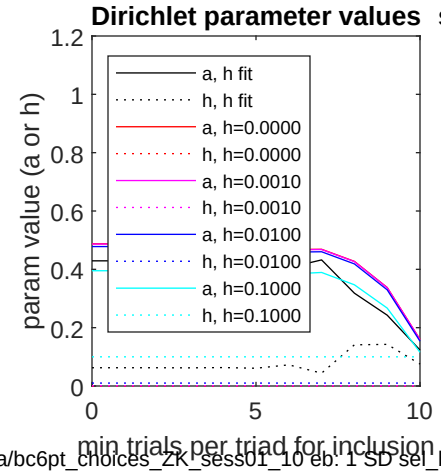

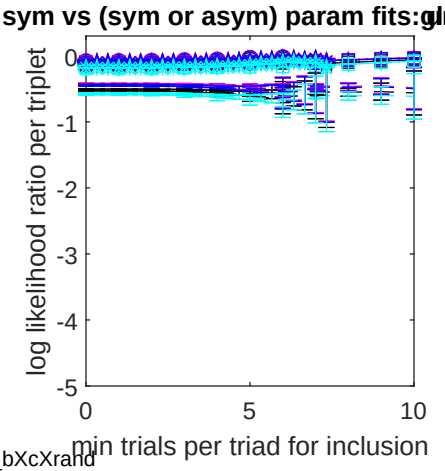

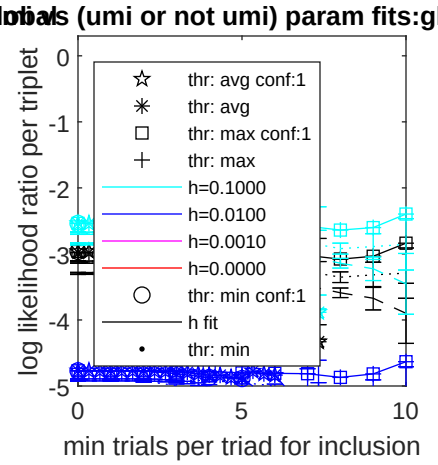

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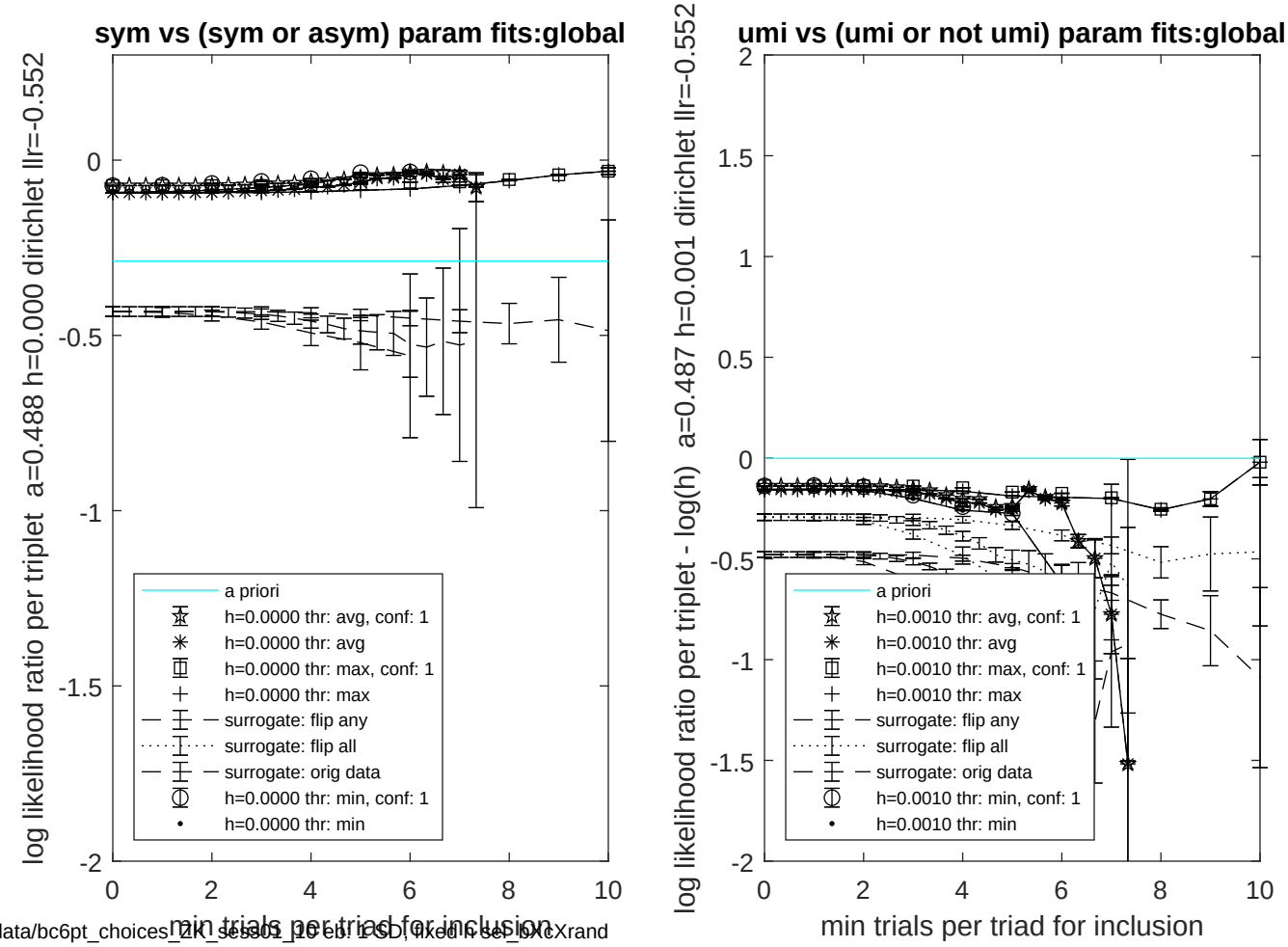

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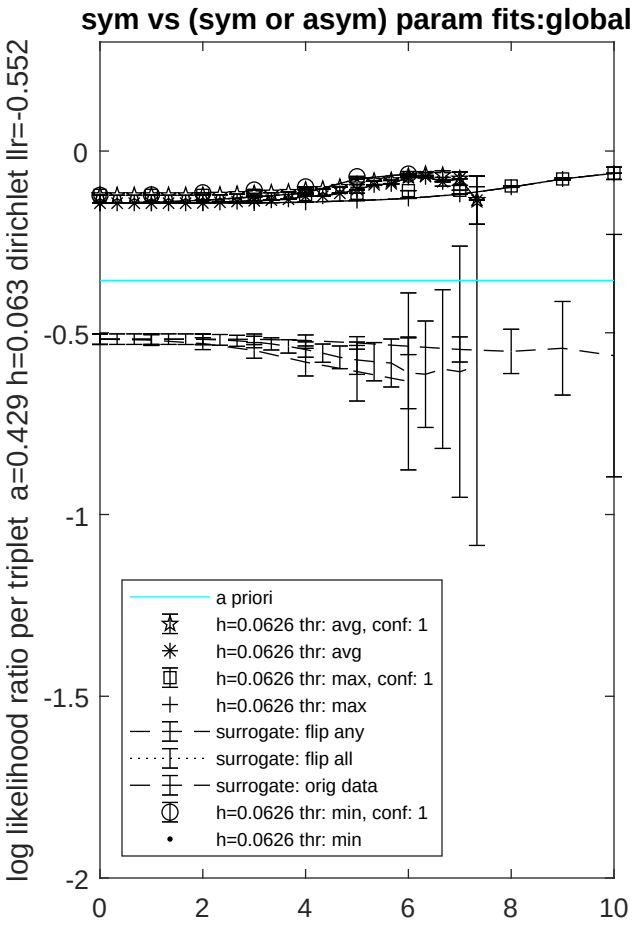

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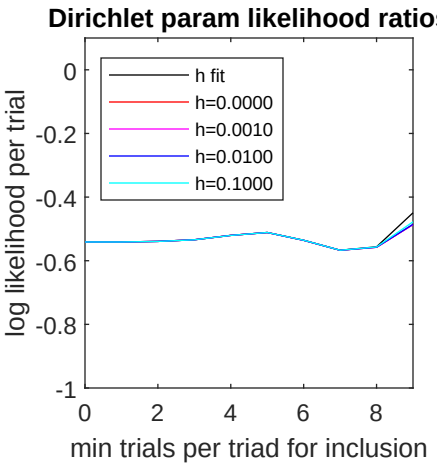

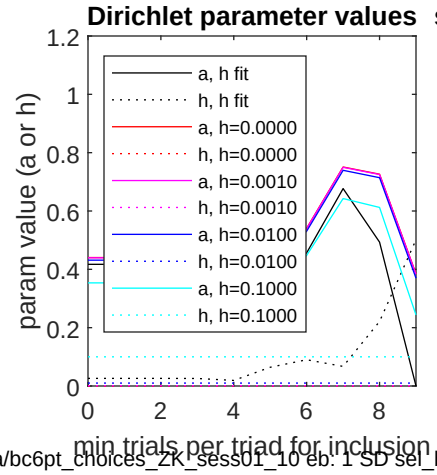

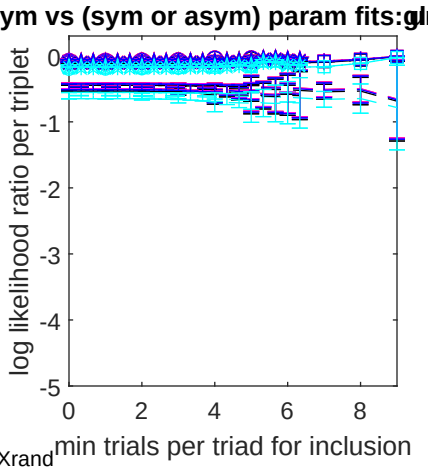

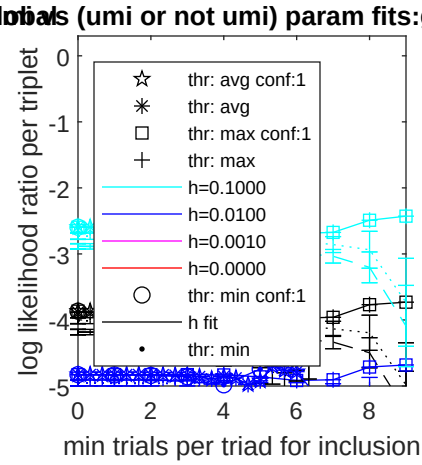

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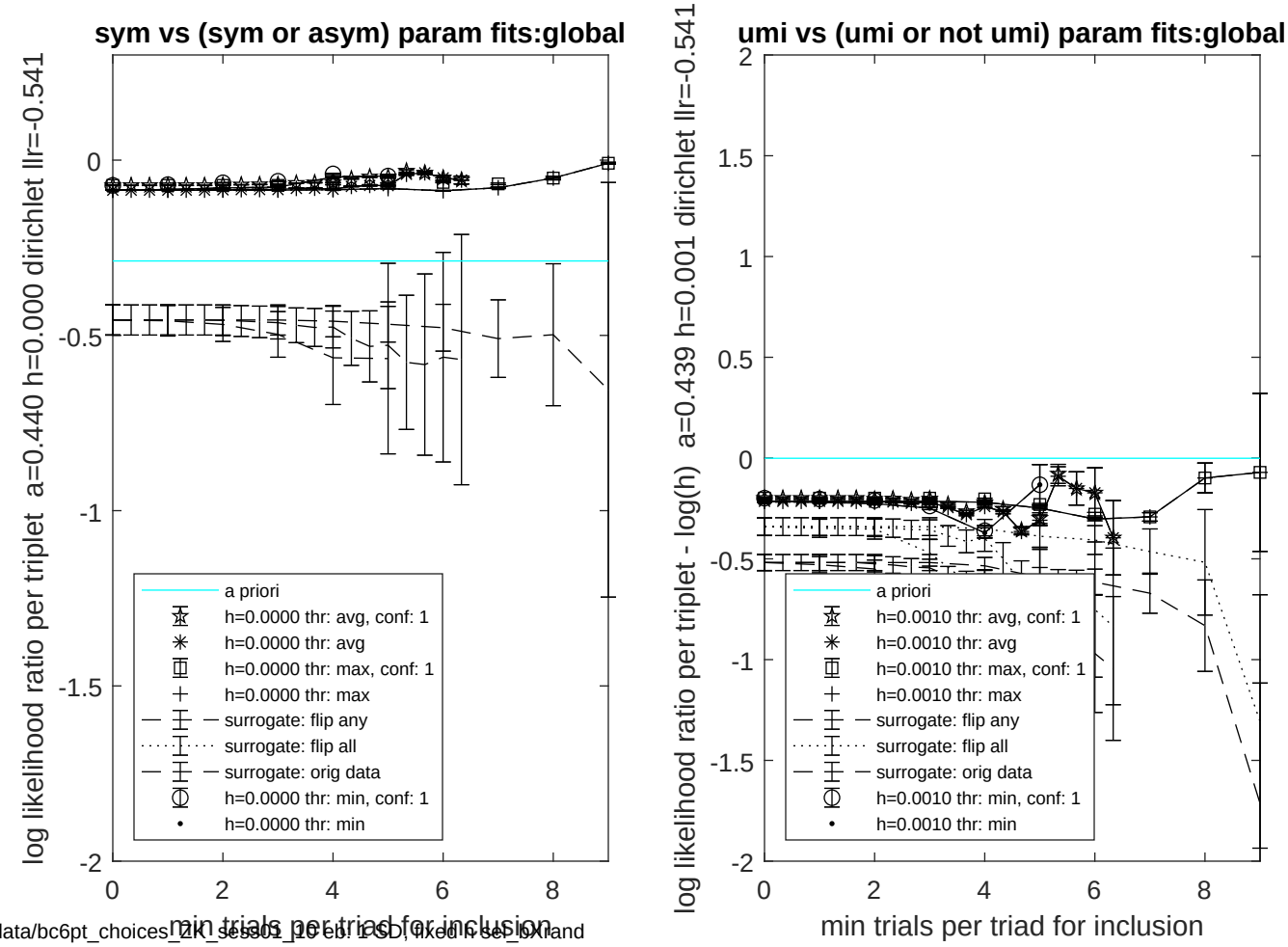

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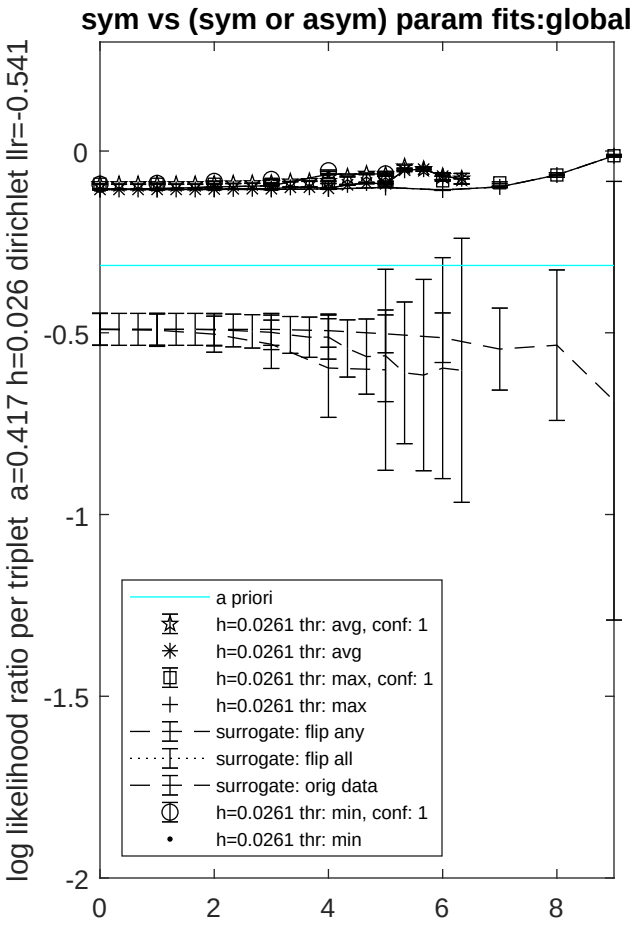

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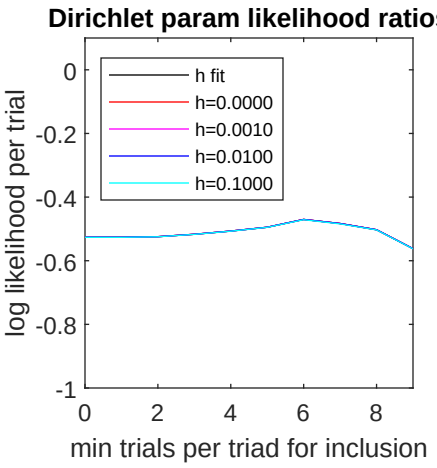

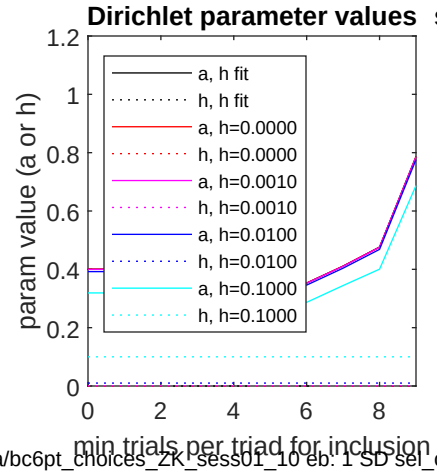

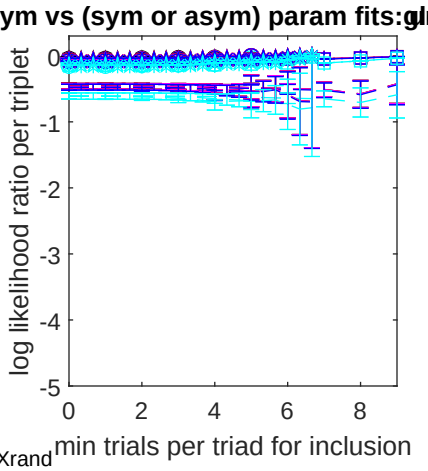

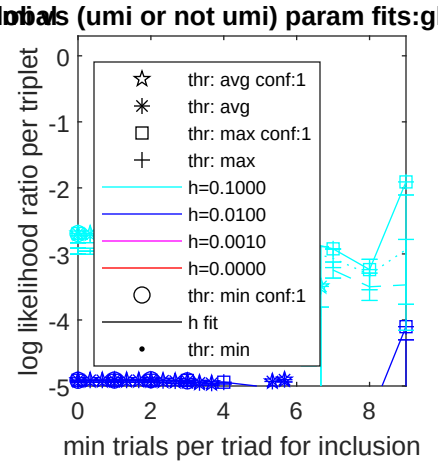

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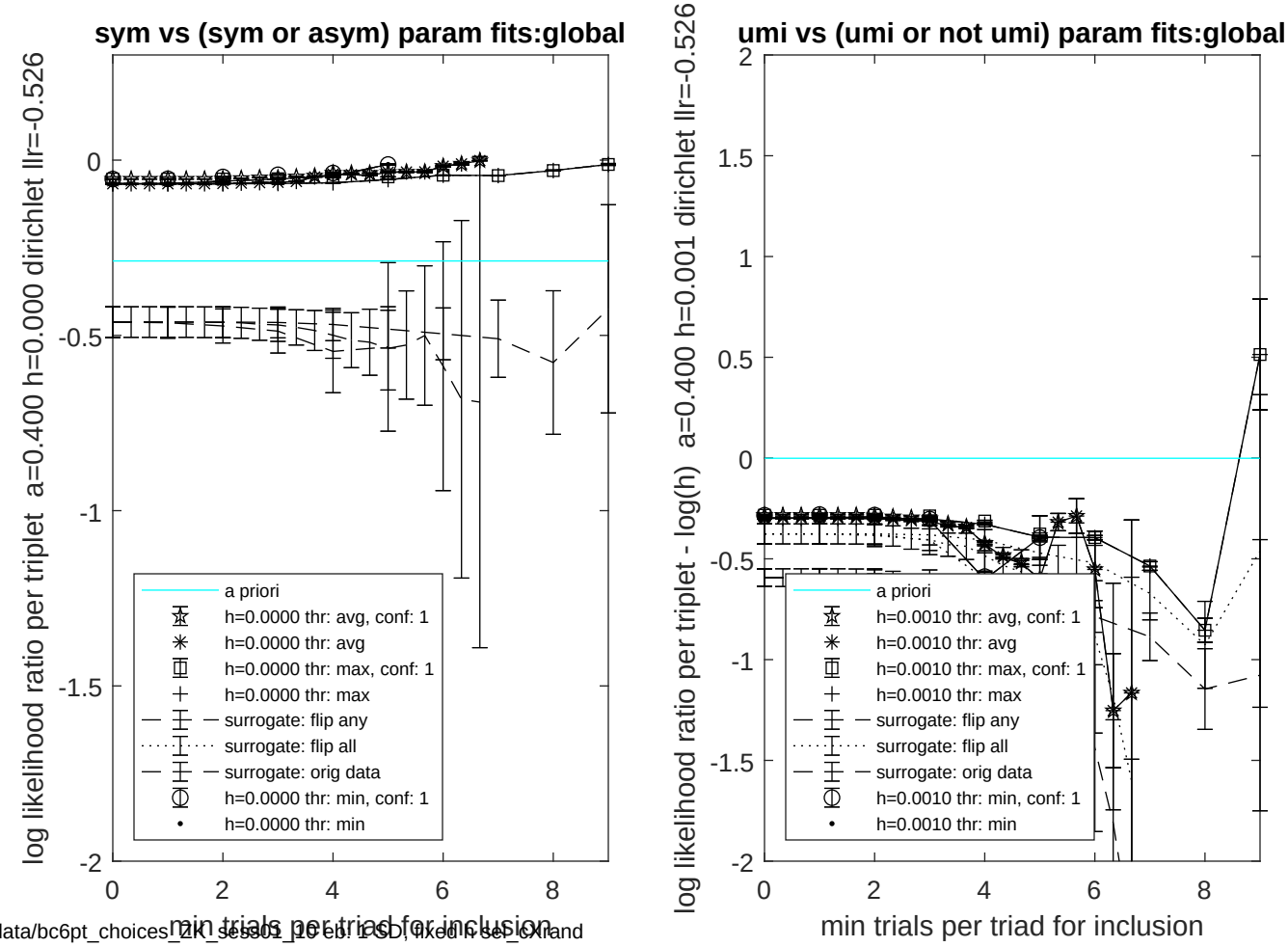

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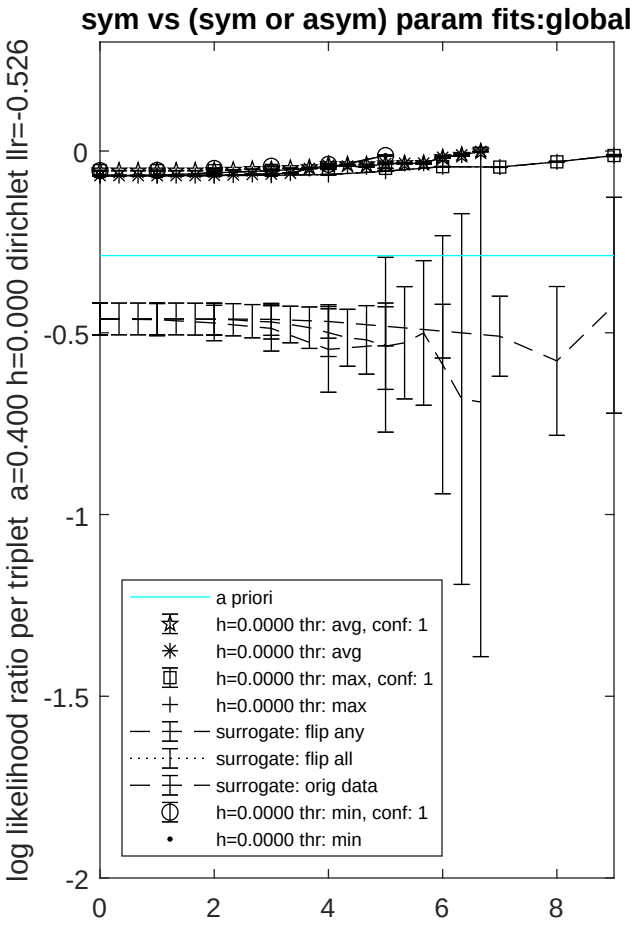

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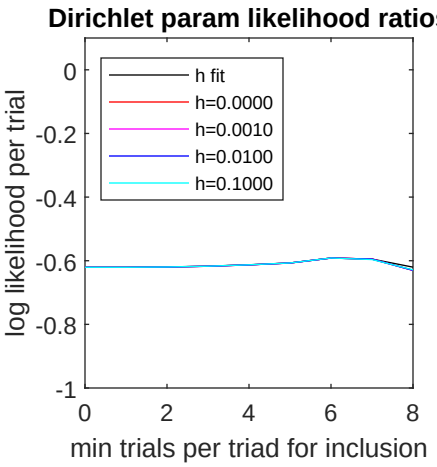

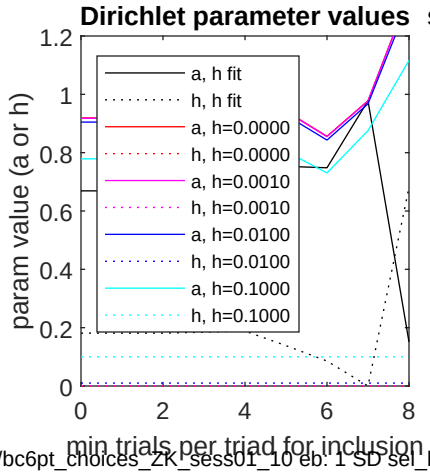

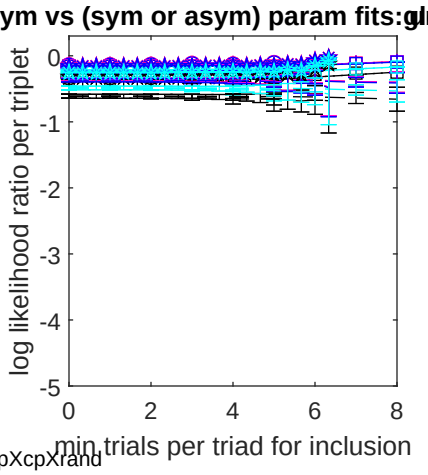

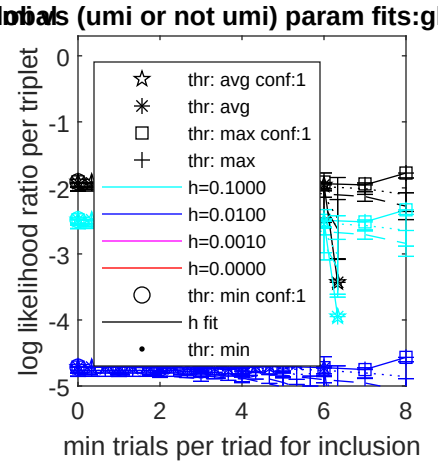

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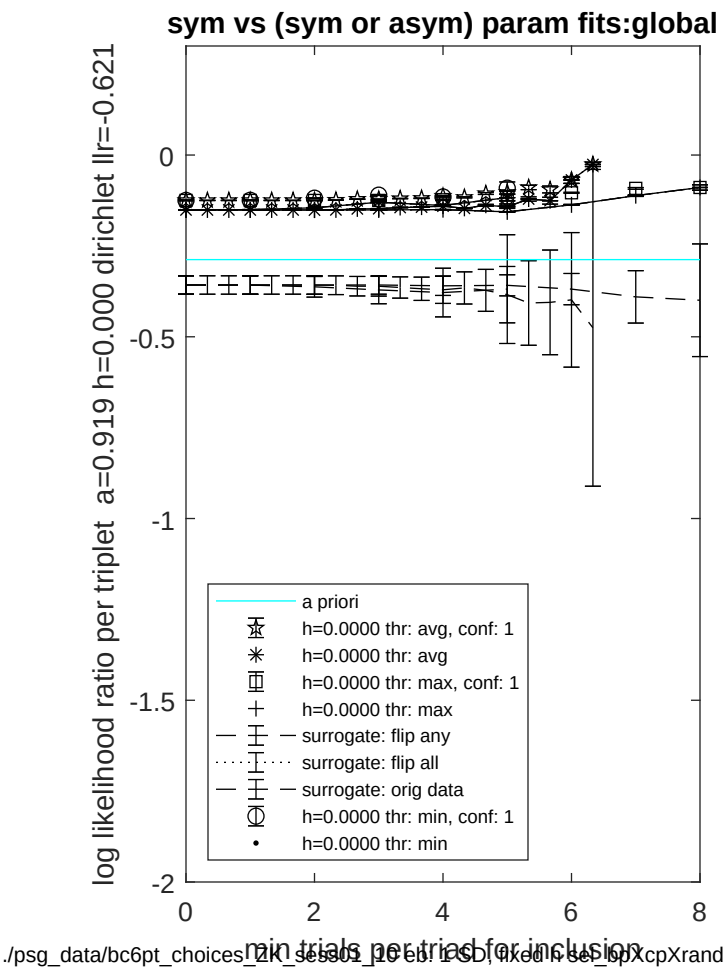
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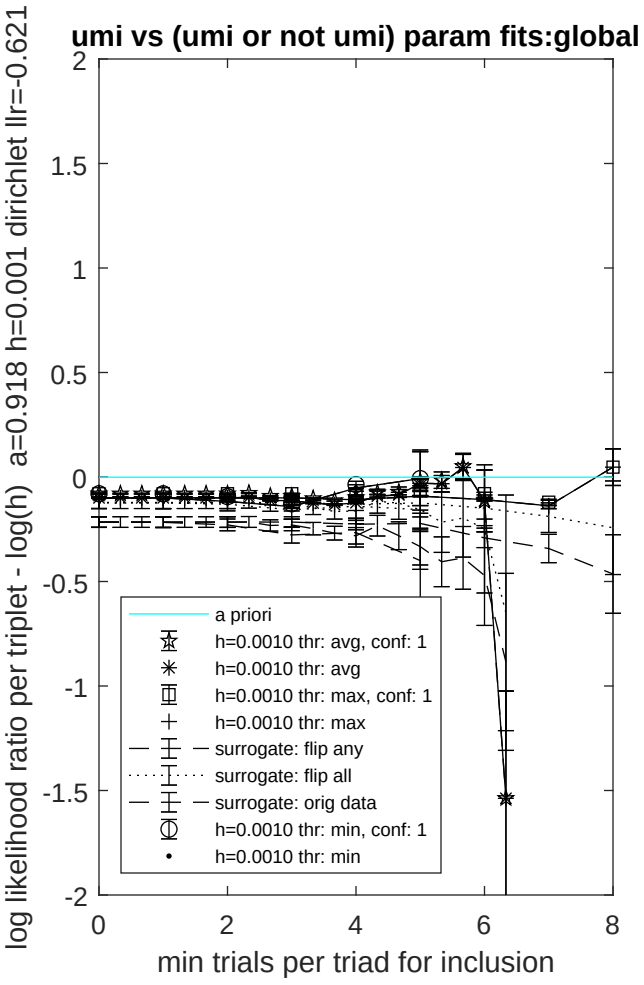
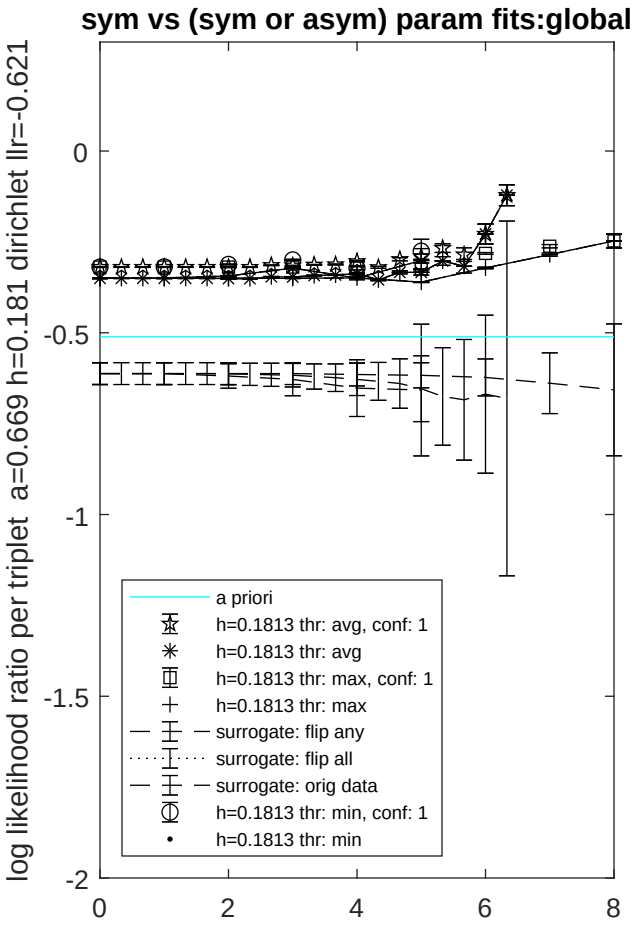
log likelihood ratio per triplet a=0.919 h=0.000 dirichlet llr=-0.621
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
./psg_data/bc6pt_choices_ZK_sess01_10 eb: 1 SD, fixed h sel_bpXcpXrand


umi vs (umi or not umi) param fits:global
log likelihood ratio per triplet - log(h) a=0.918 h=0.001 dirichlet llr=-0.621
min trials per triad for inclusion
a priori
h=0.0010 thr: avg, conf: 1
h=0.0010 thr: avg
h=0.0010 thr: max, conf: 1
h=0.0010 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0010 thr: min, conf: 1
h=0.0010 thr: min



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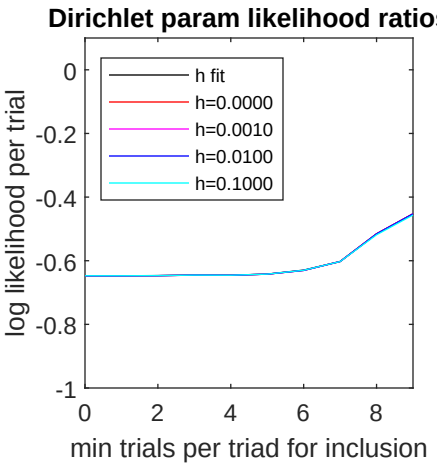

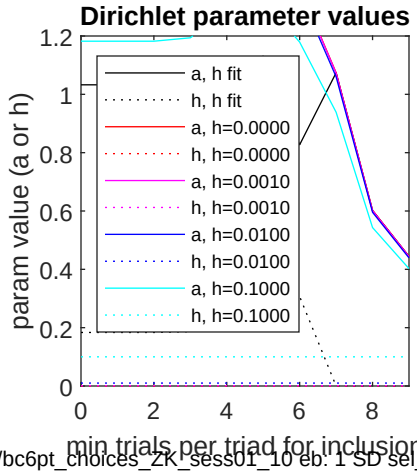

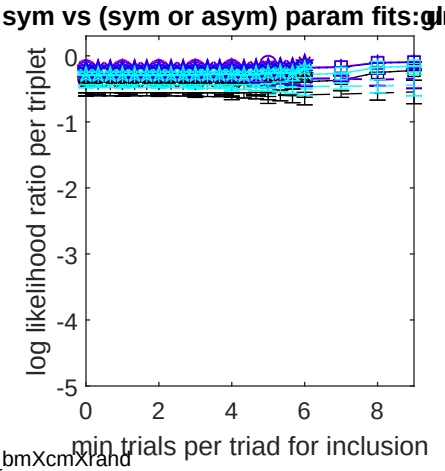

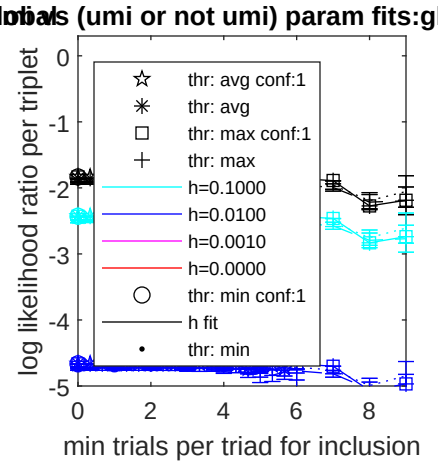

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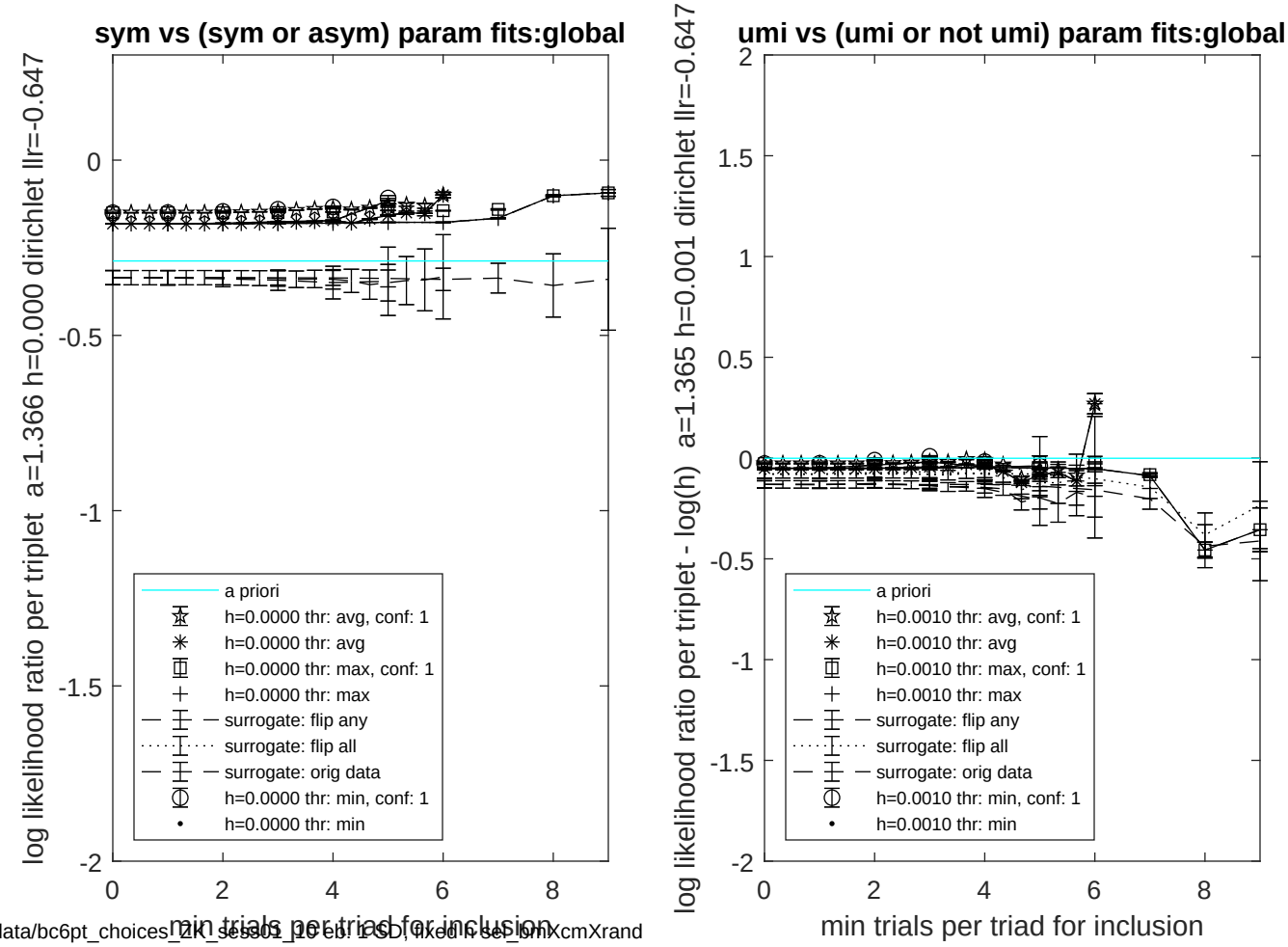

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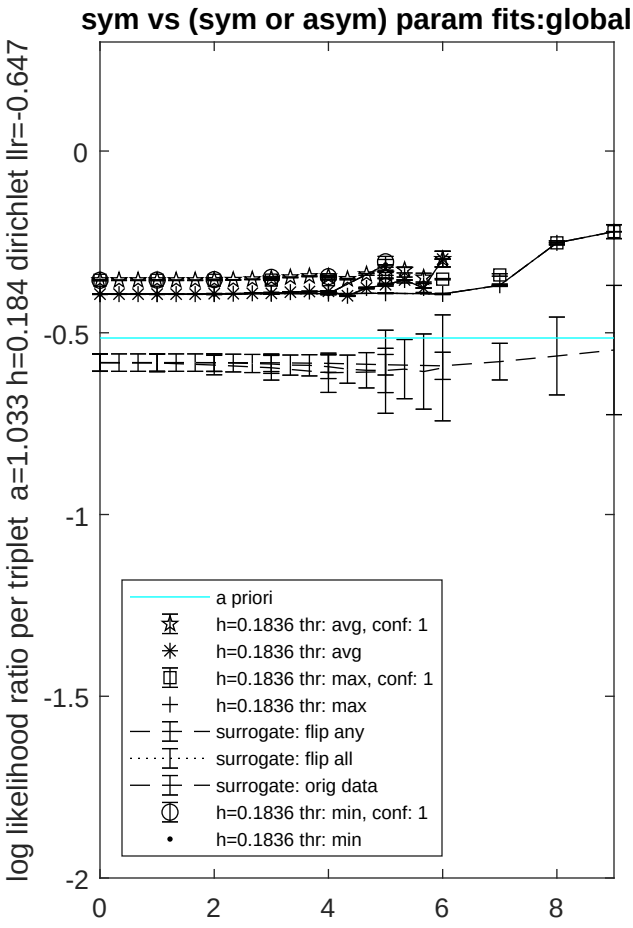

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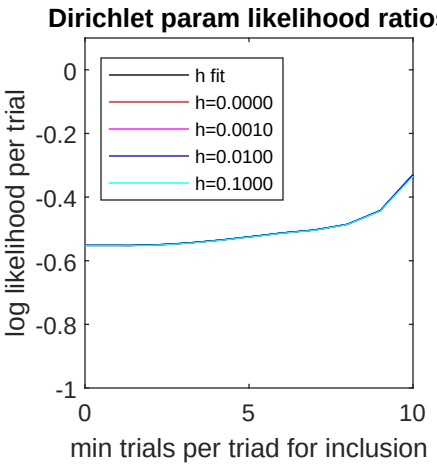

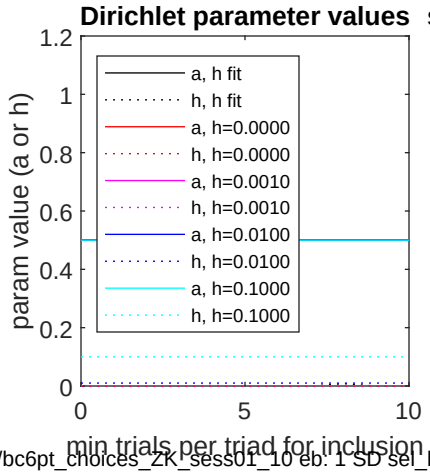

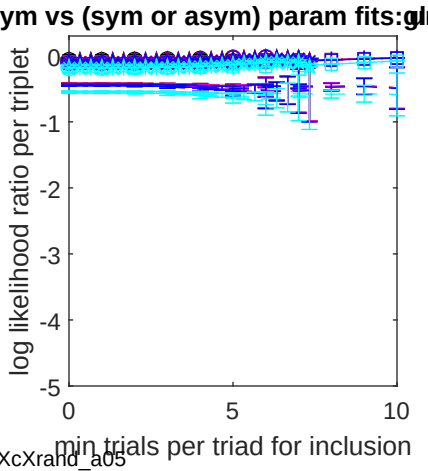

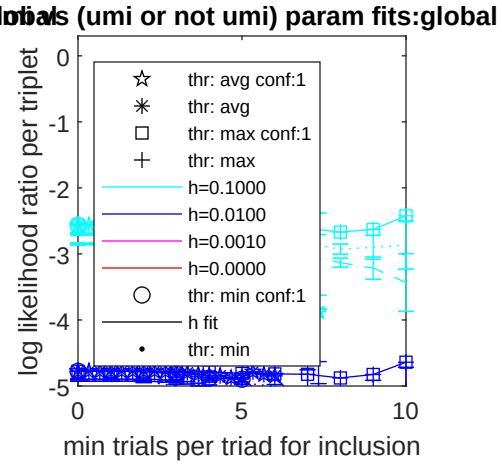

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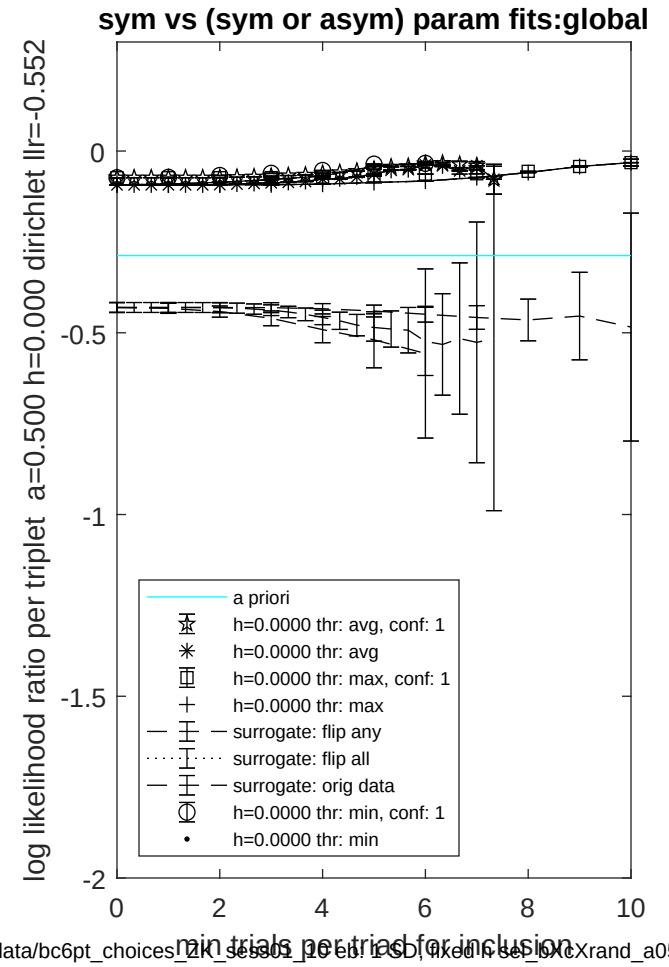

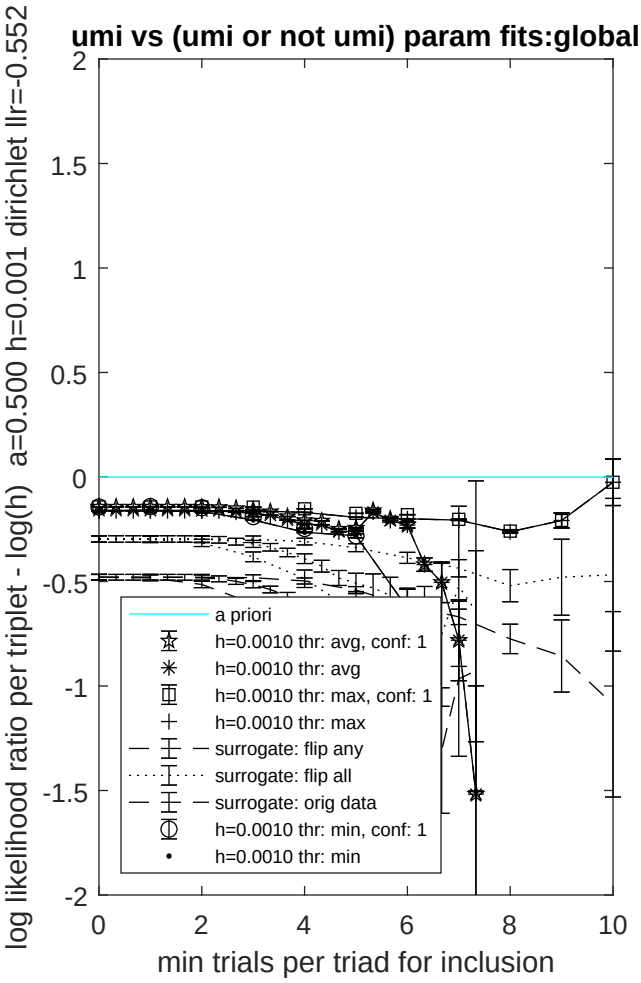

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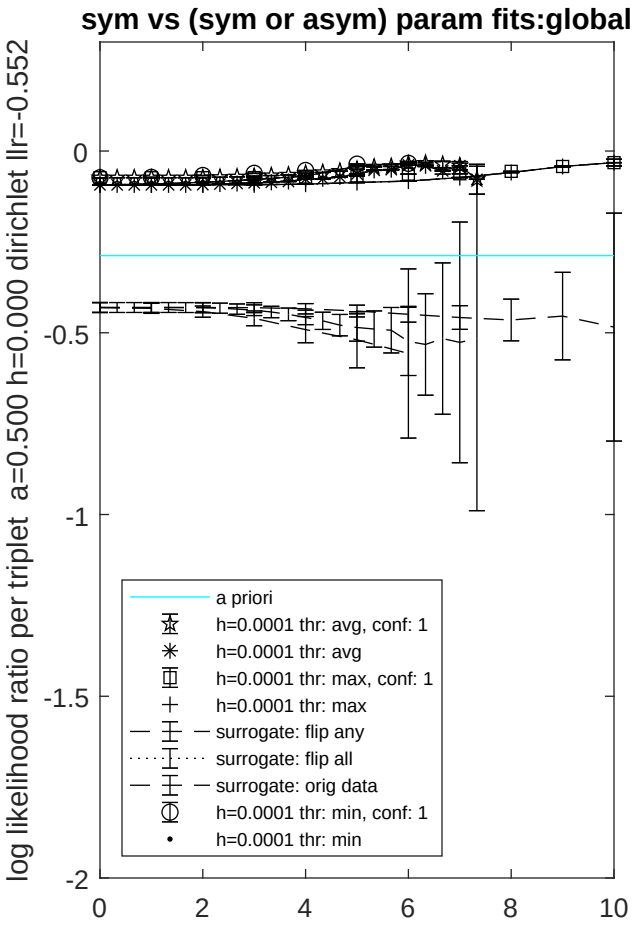

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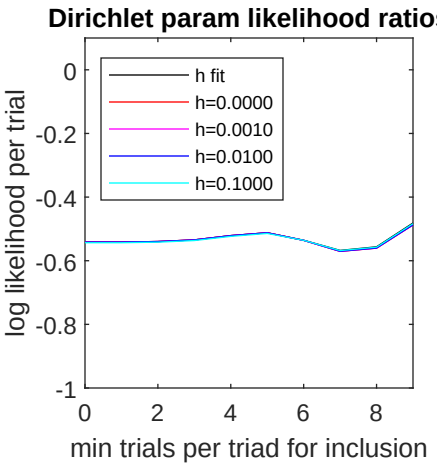

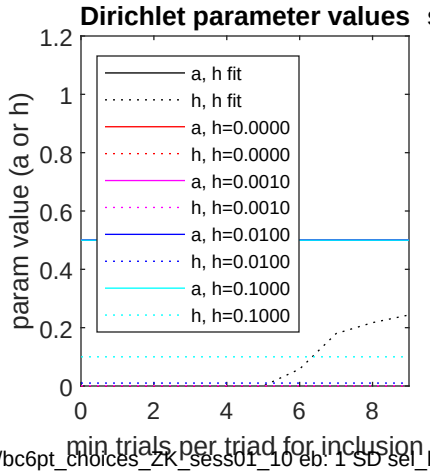

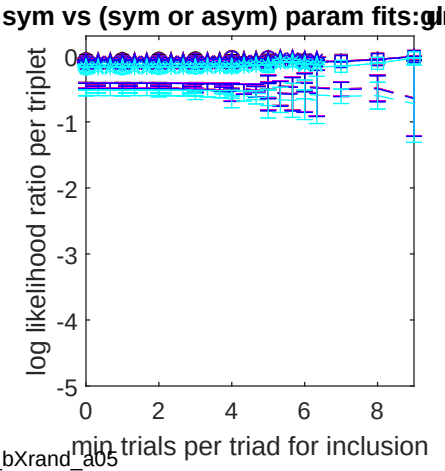

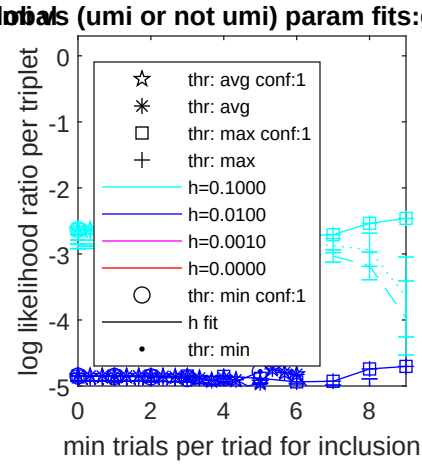

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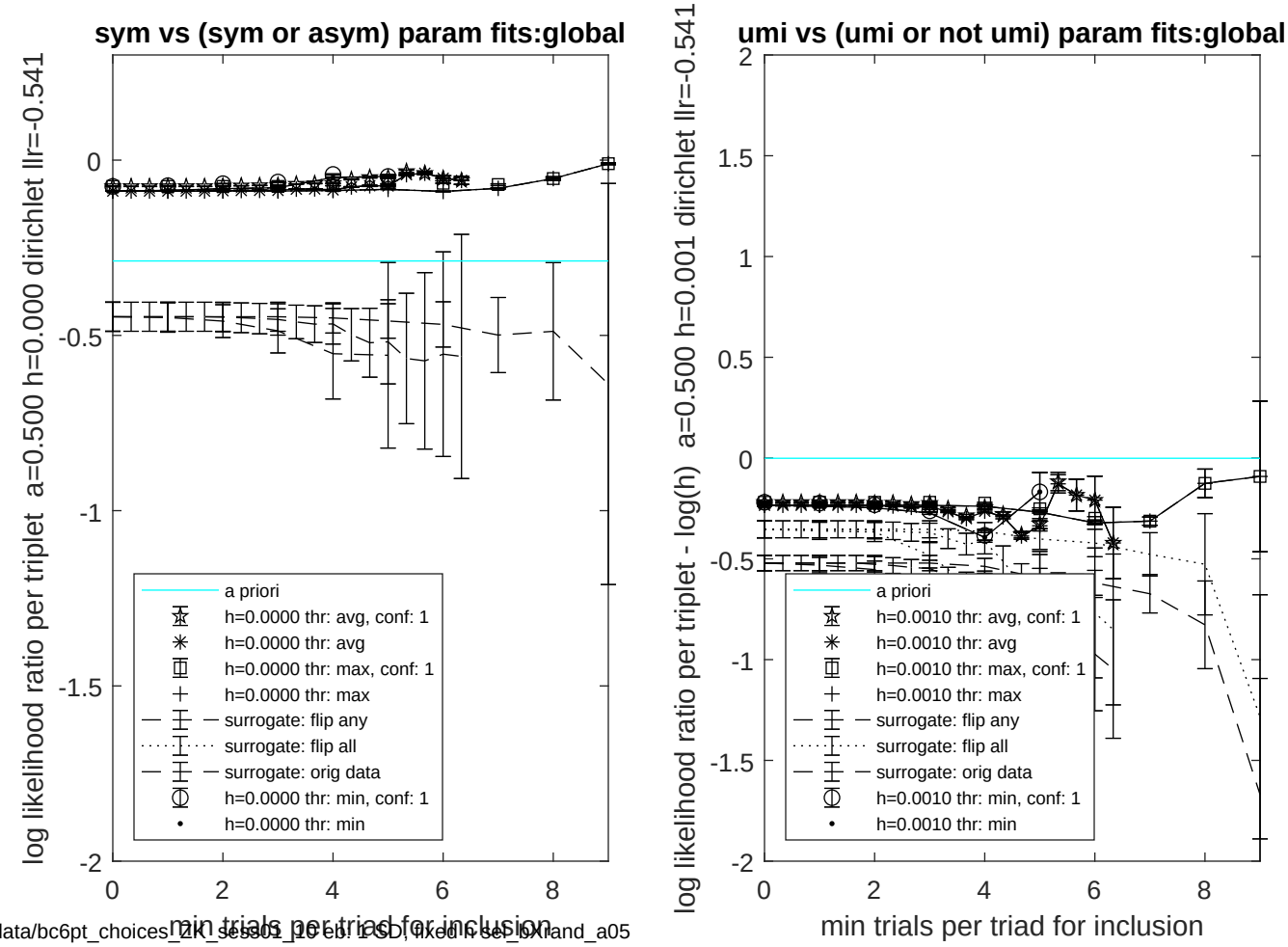

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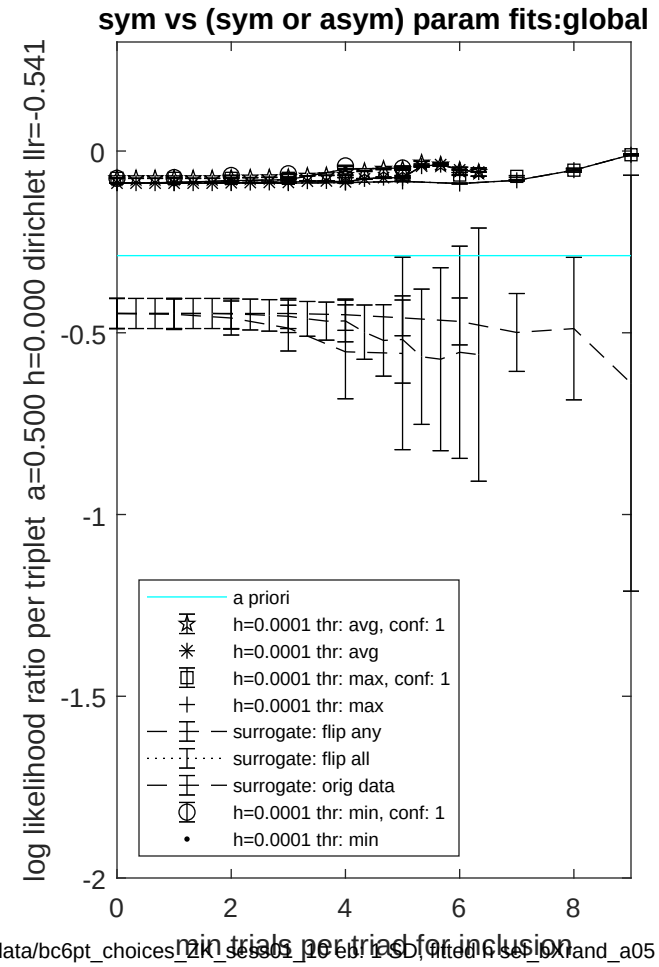

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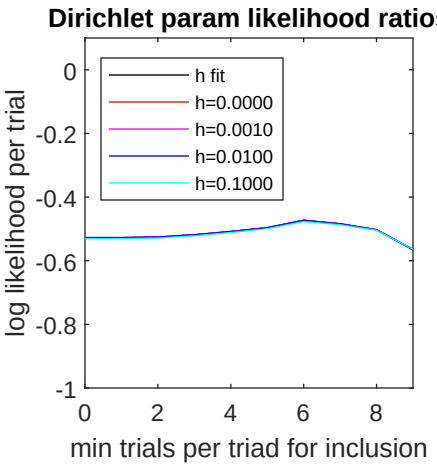

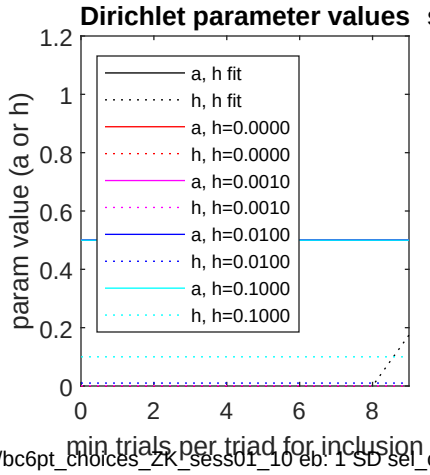

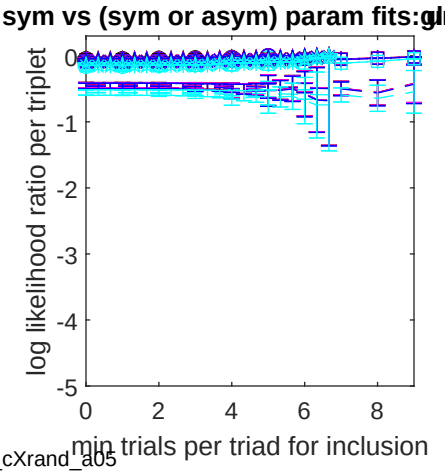

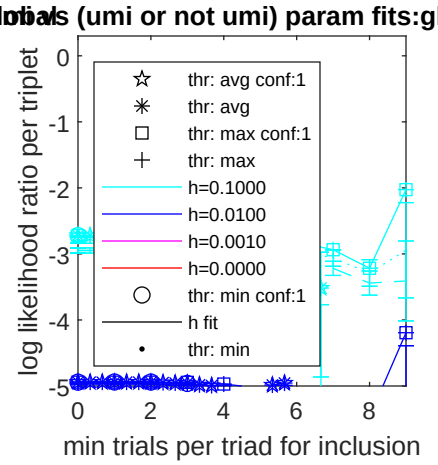

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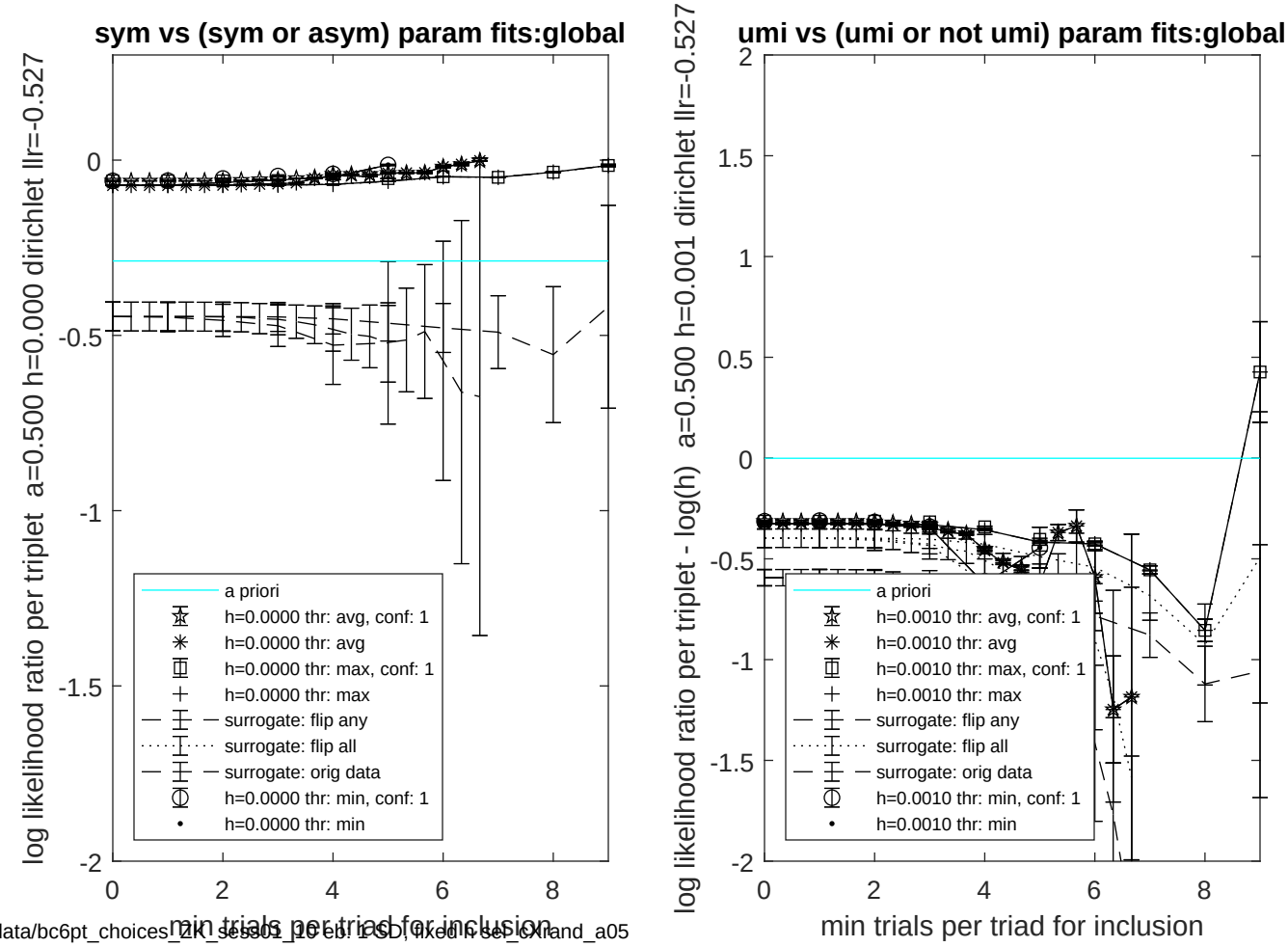

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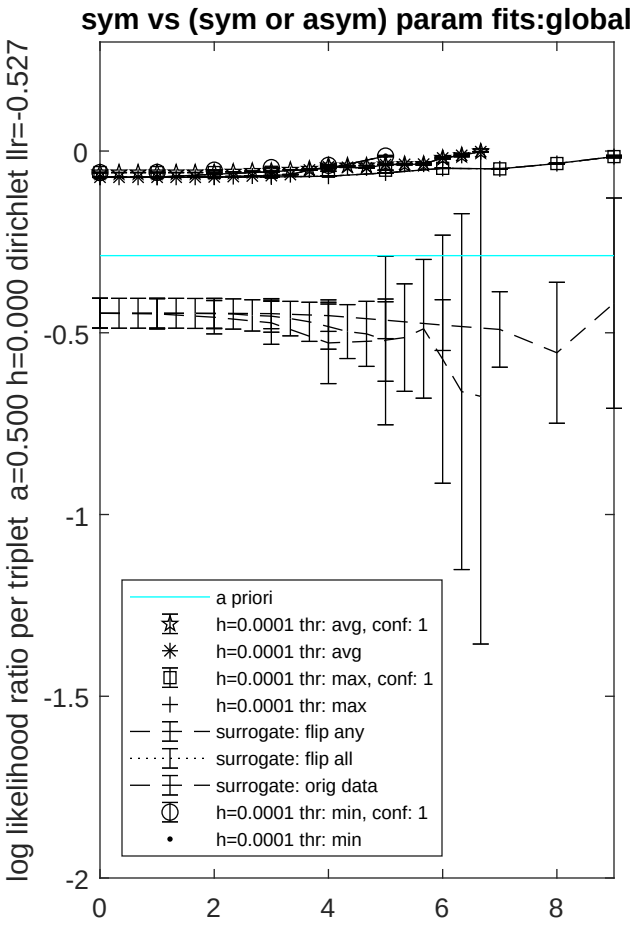

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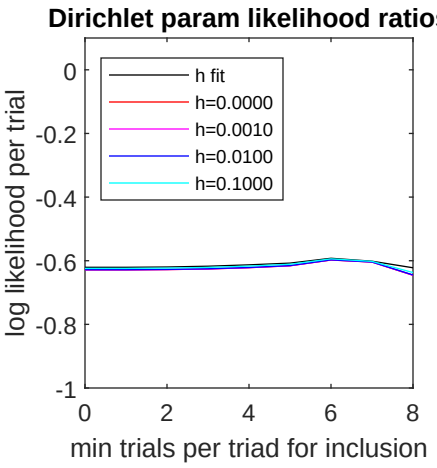

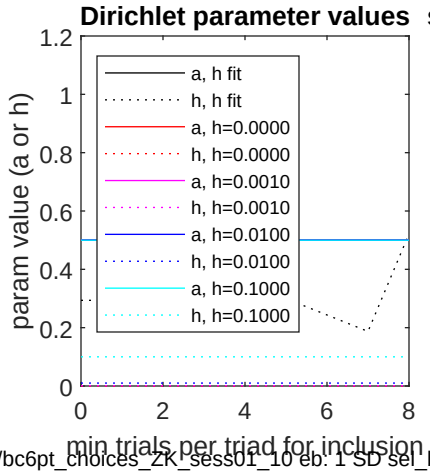

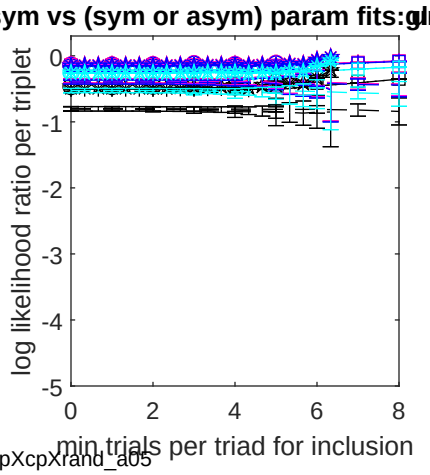

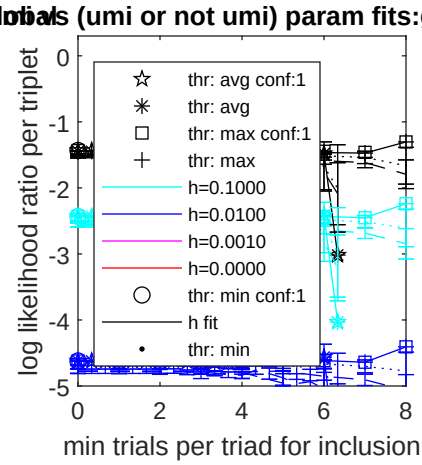

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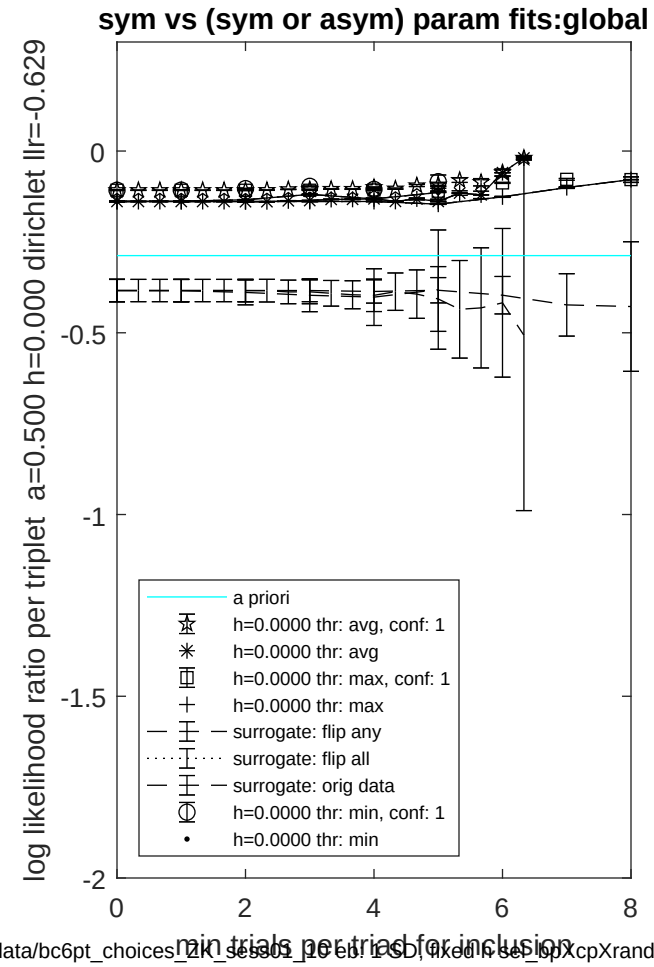

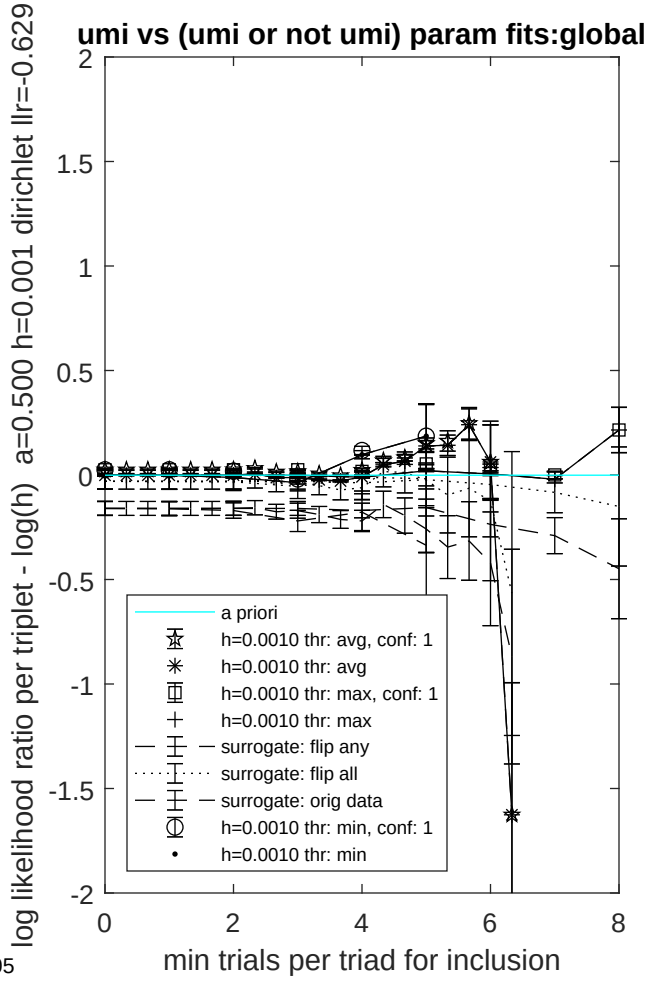

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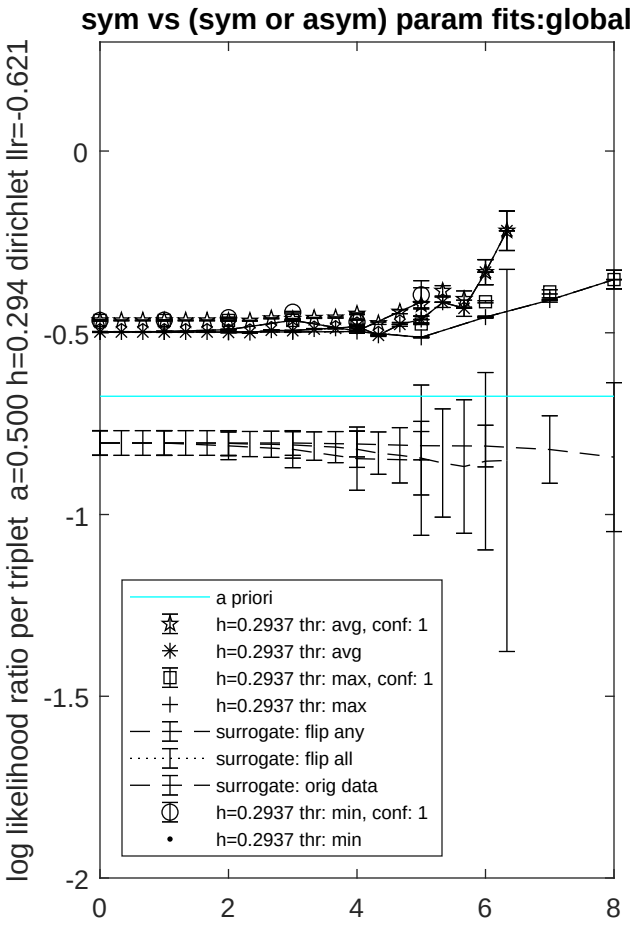

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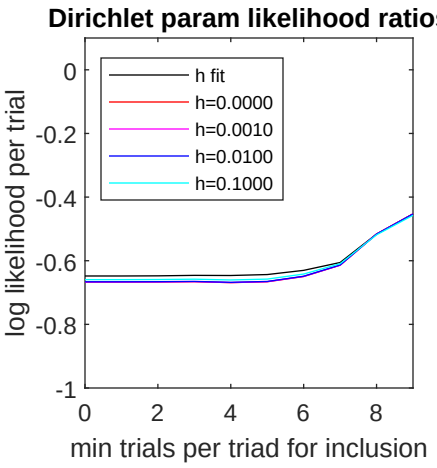

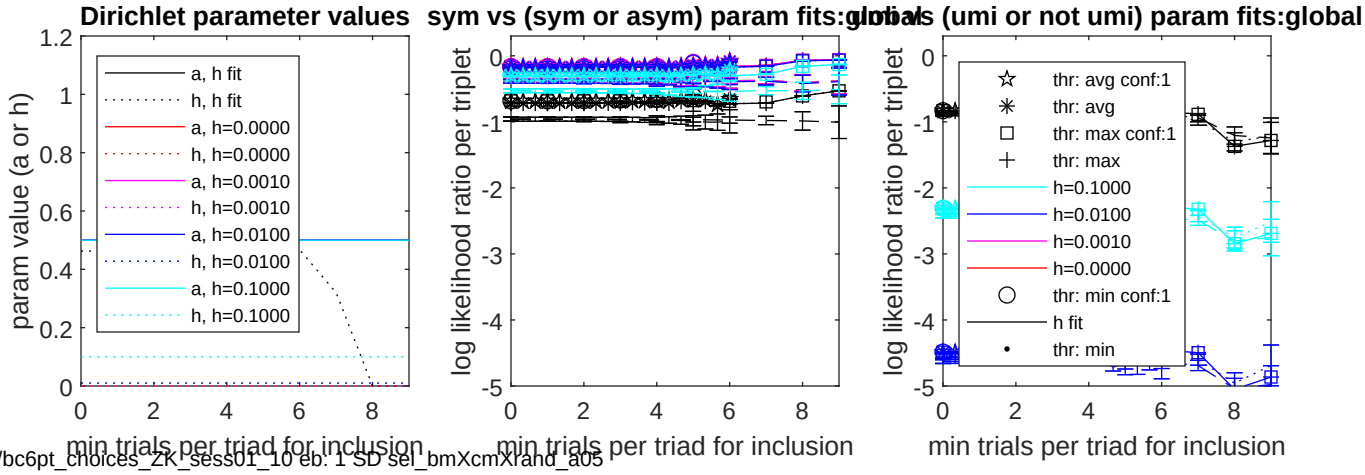

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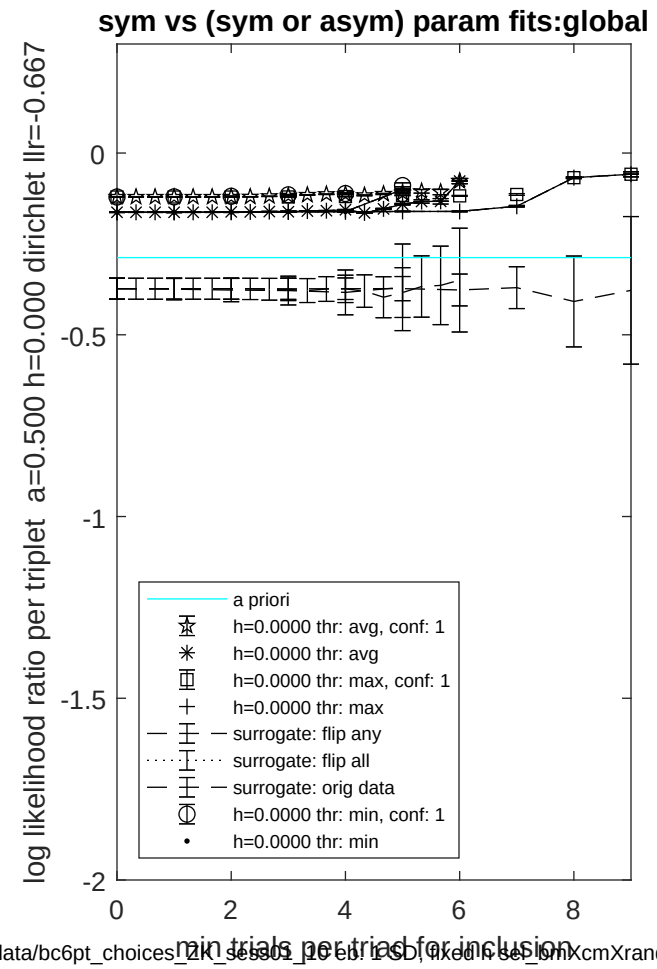

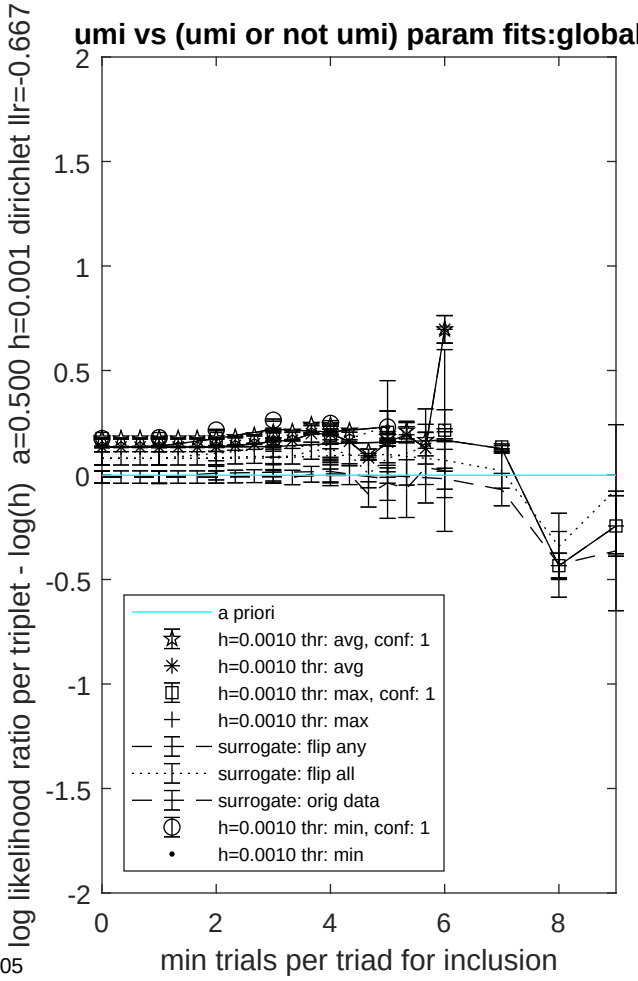

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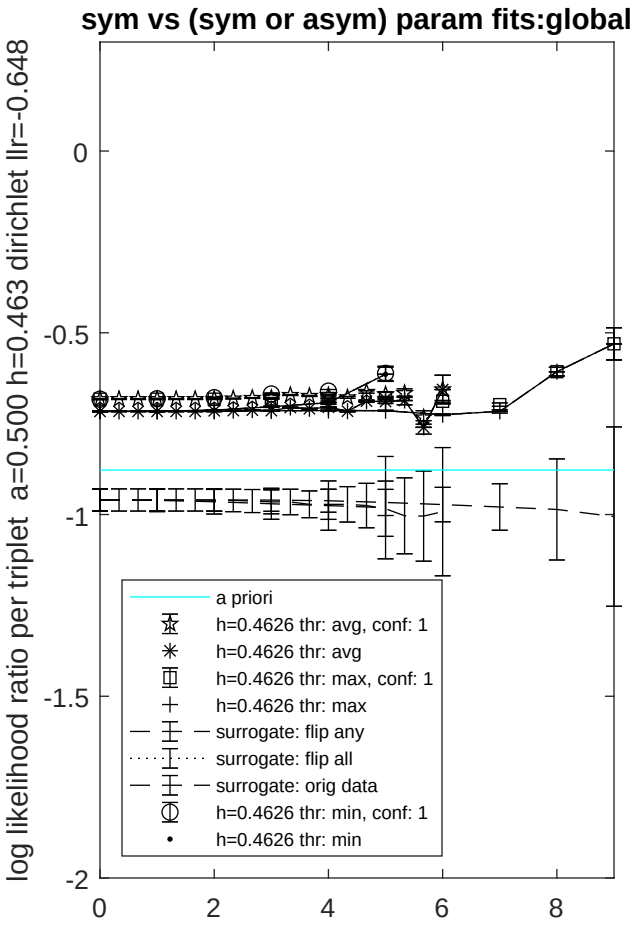

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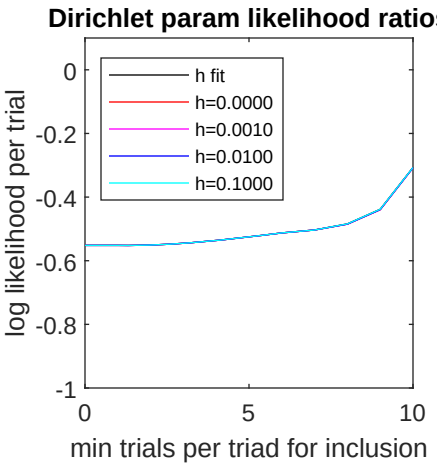

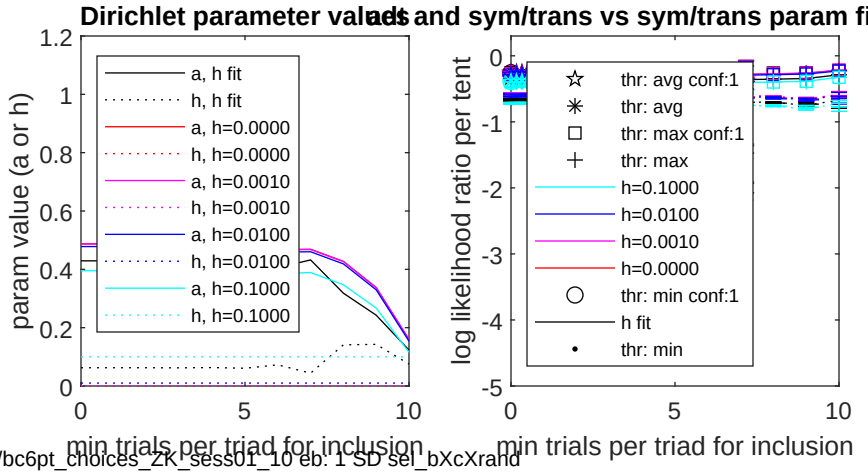

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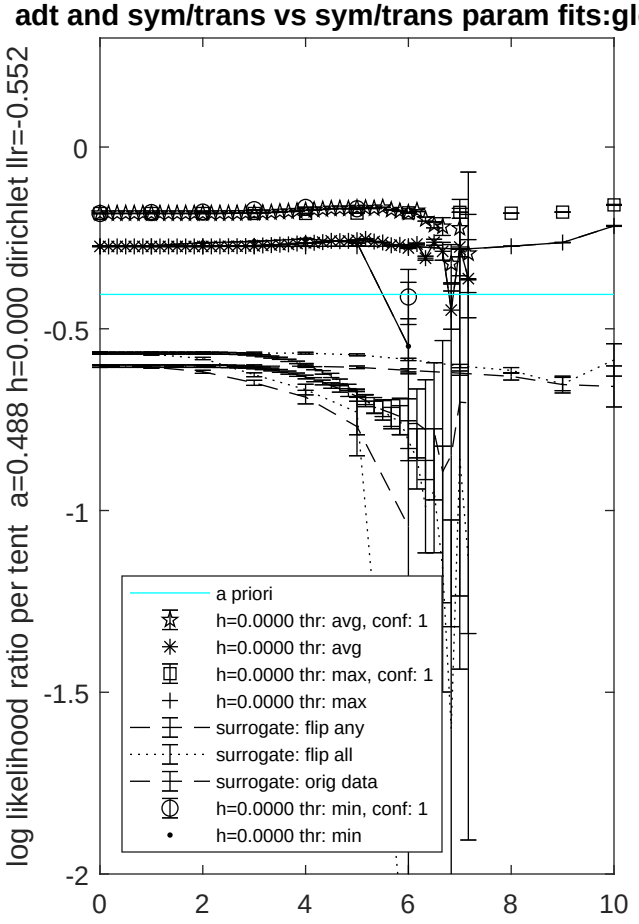

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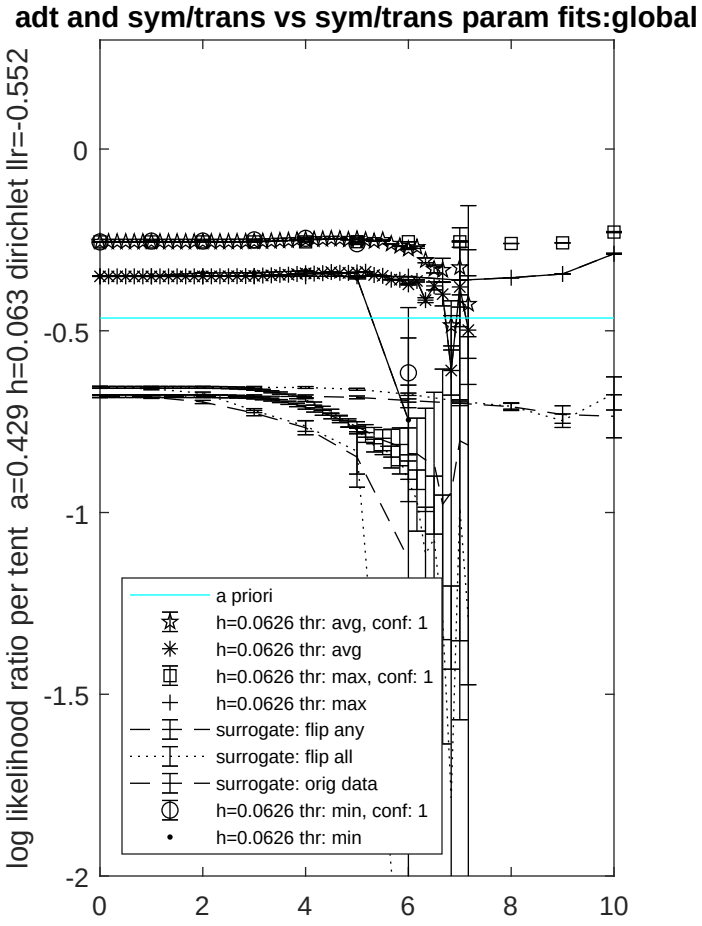

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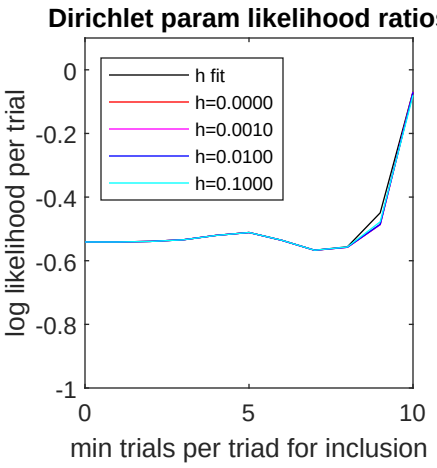

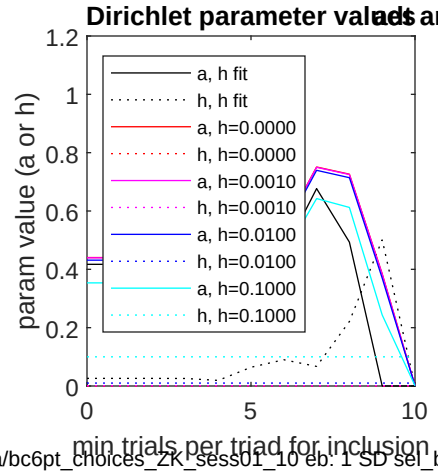

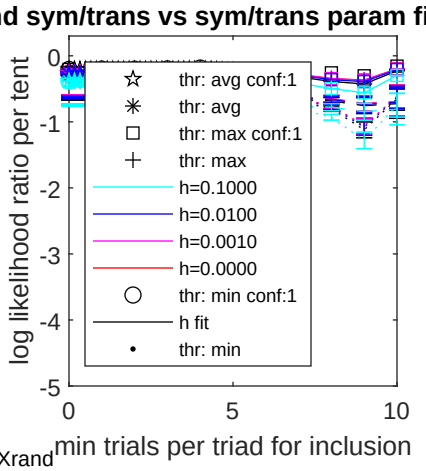

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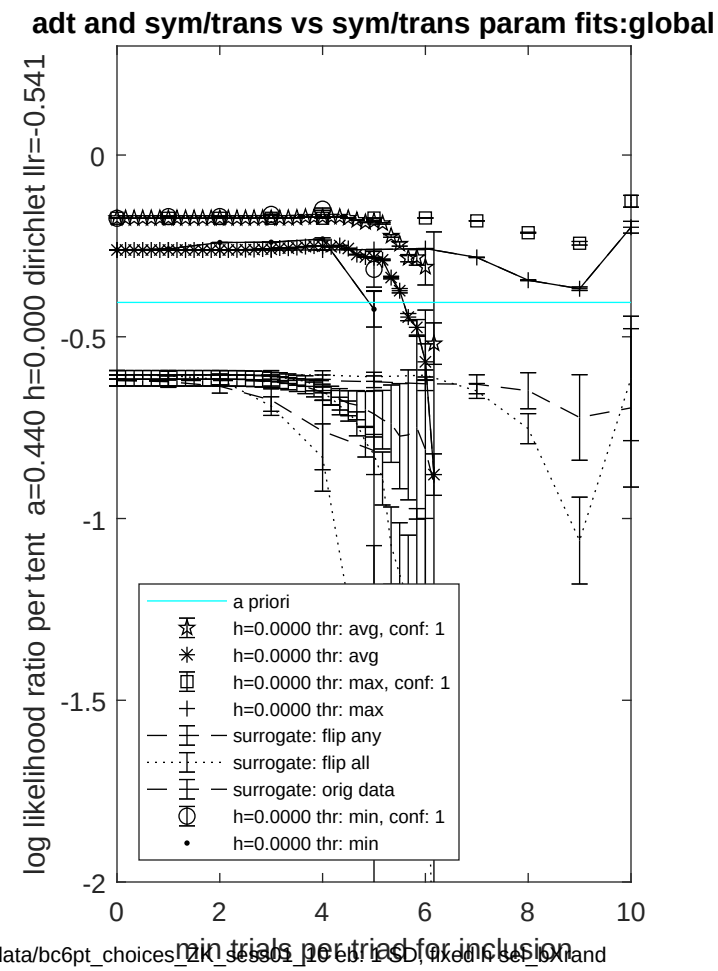

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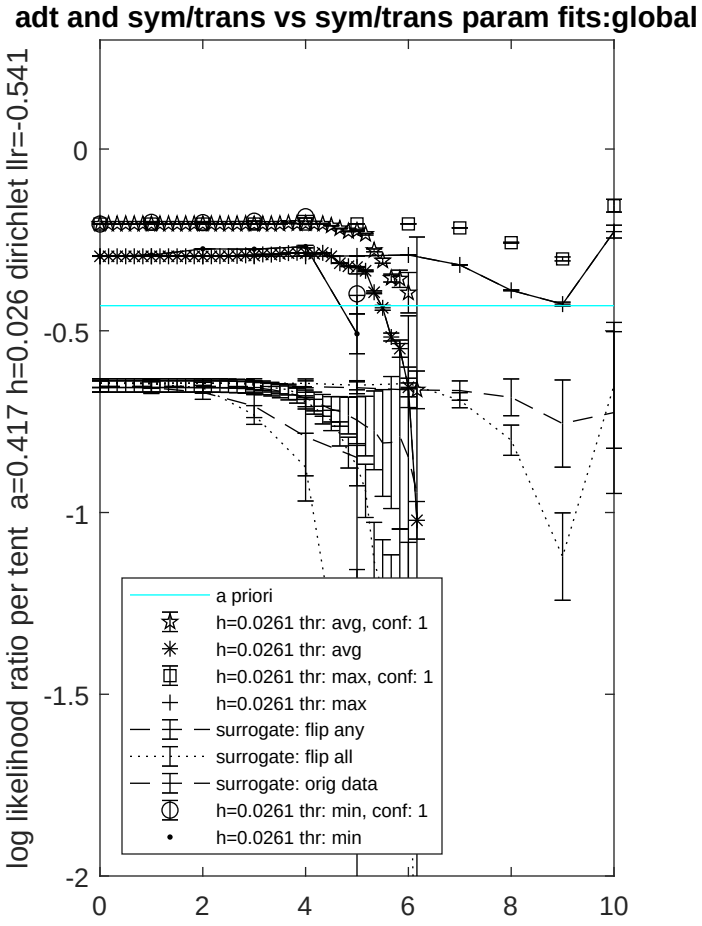

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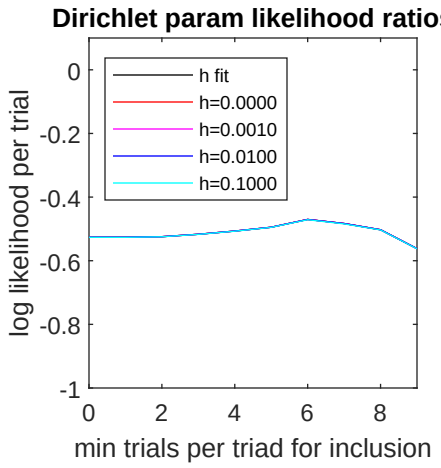

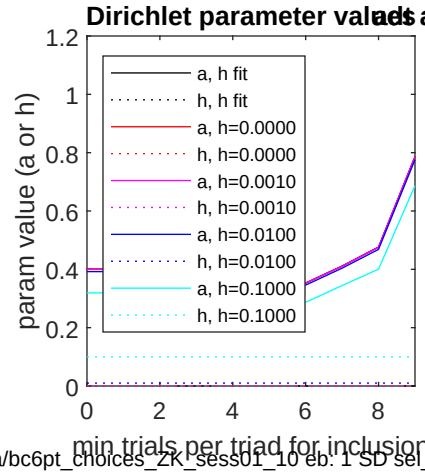

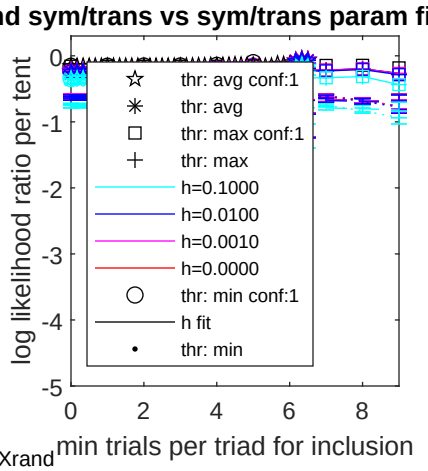

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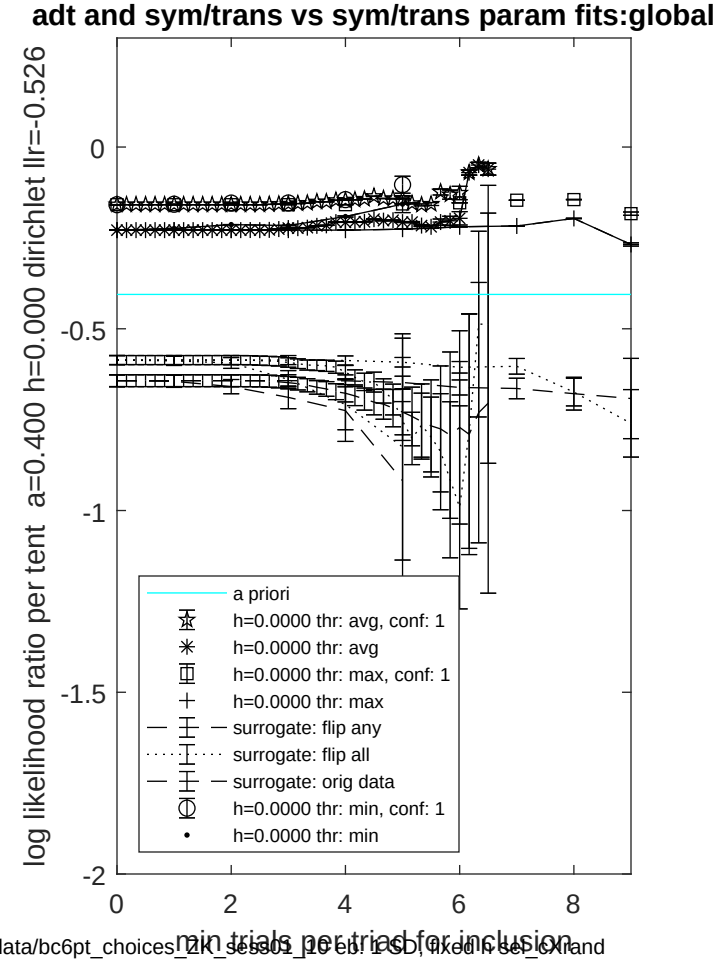

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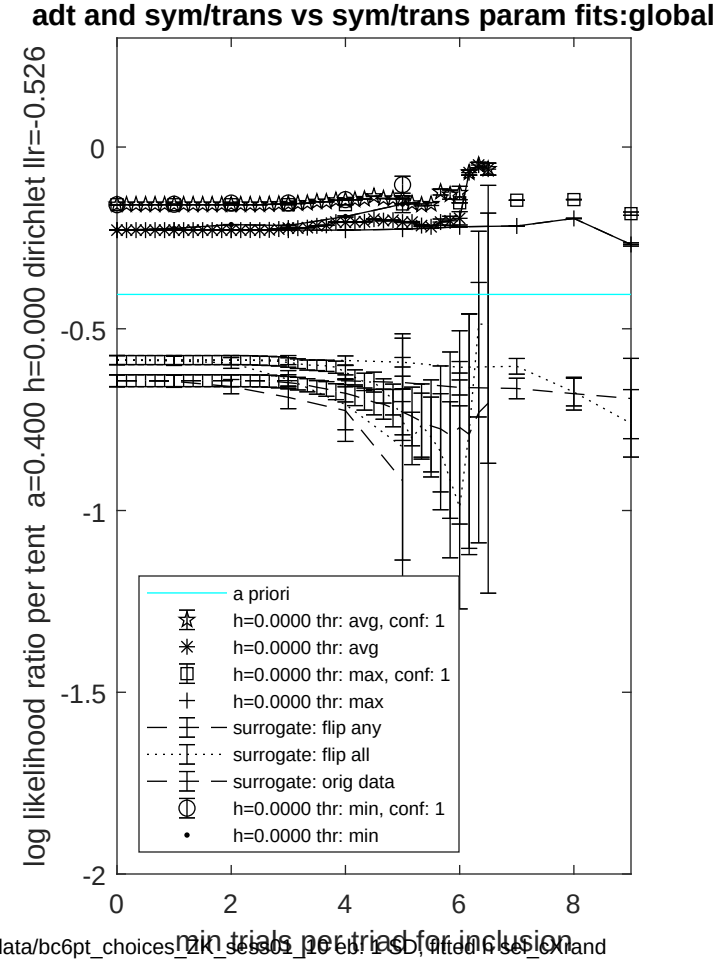

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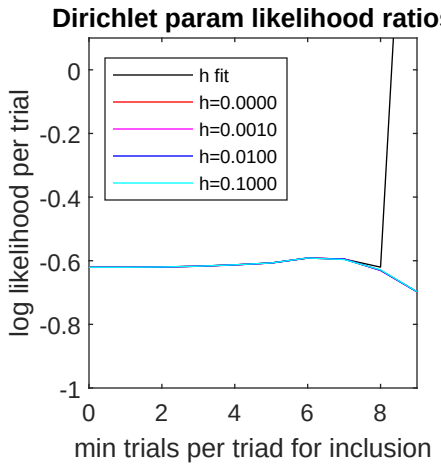

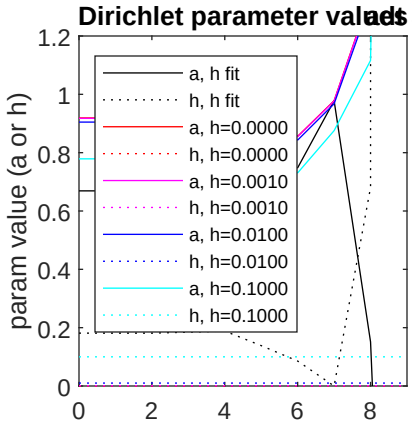

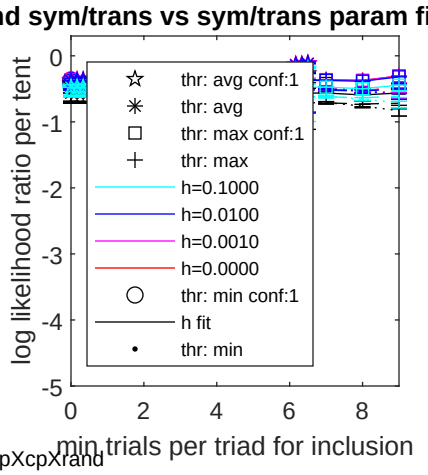

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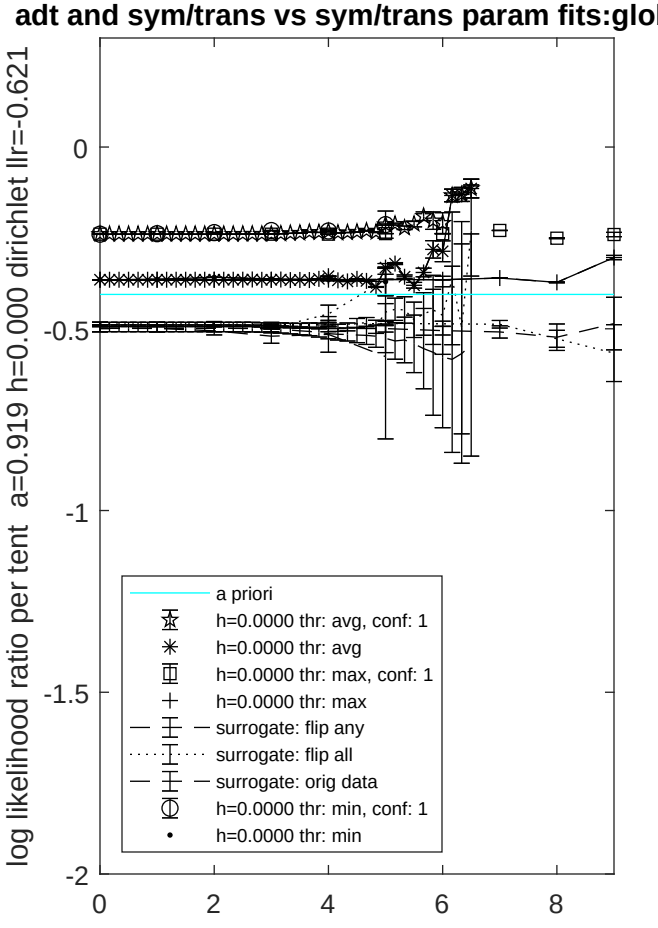

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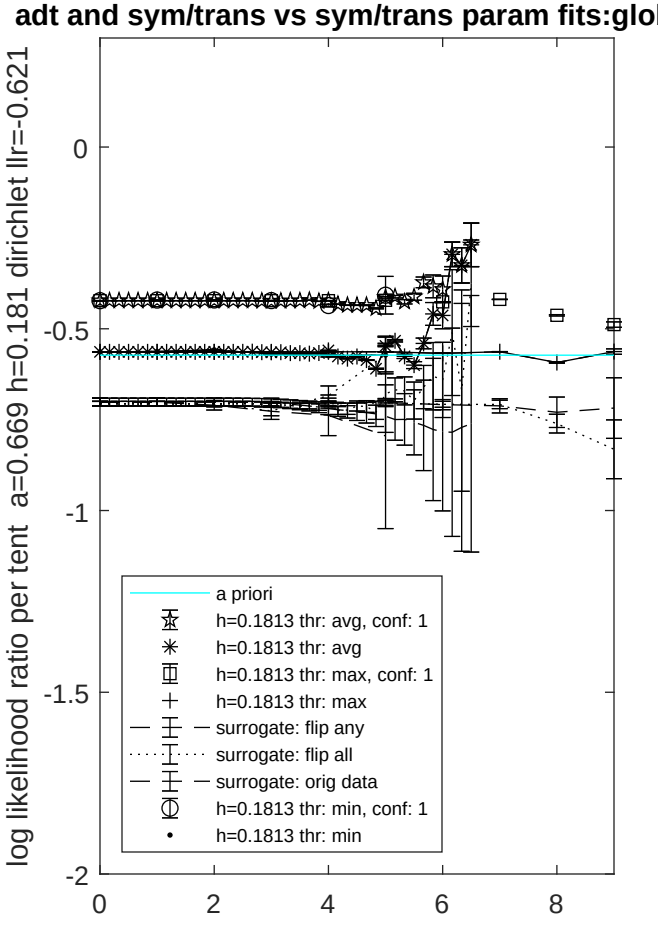

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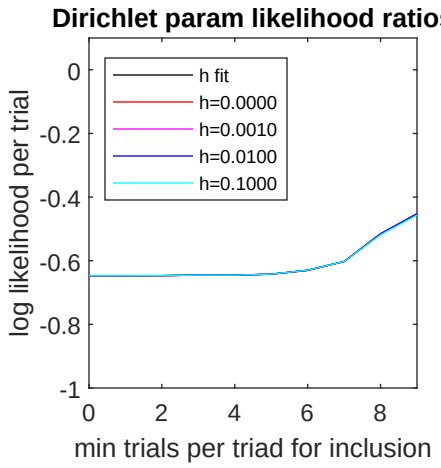

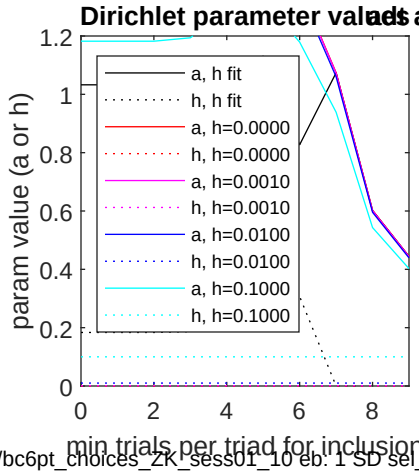

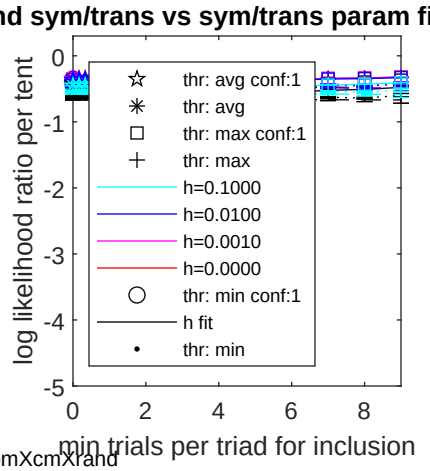

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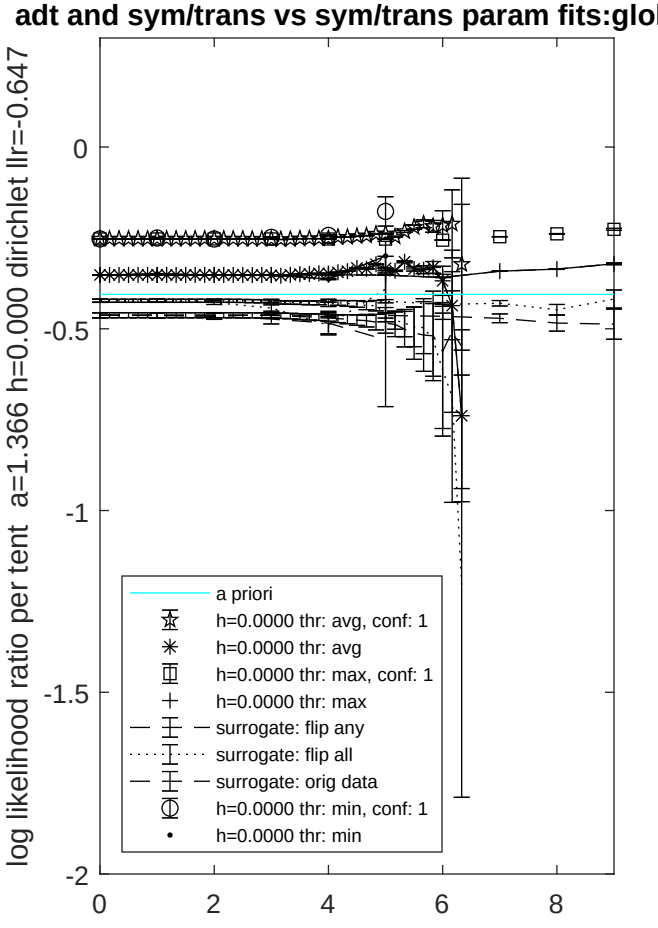

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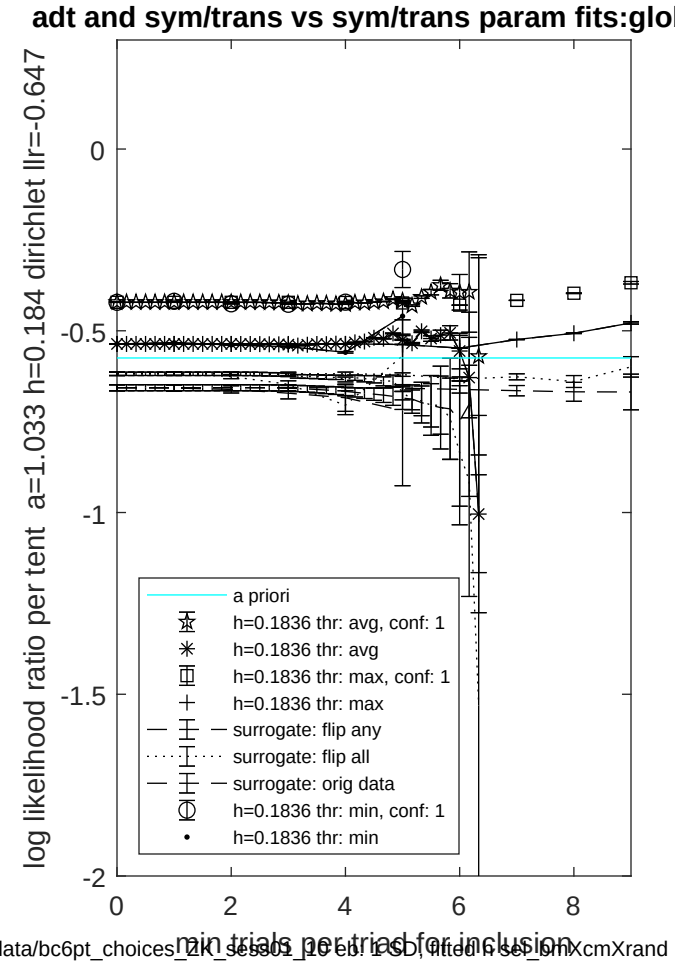

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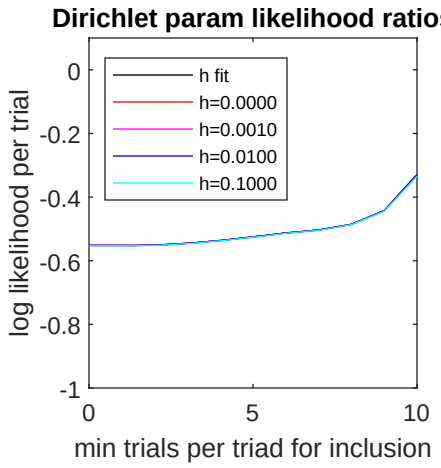

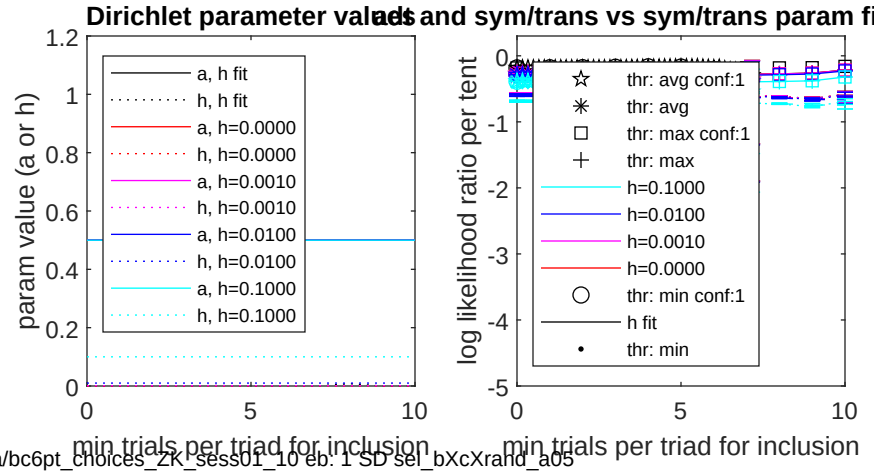

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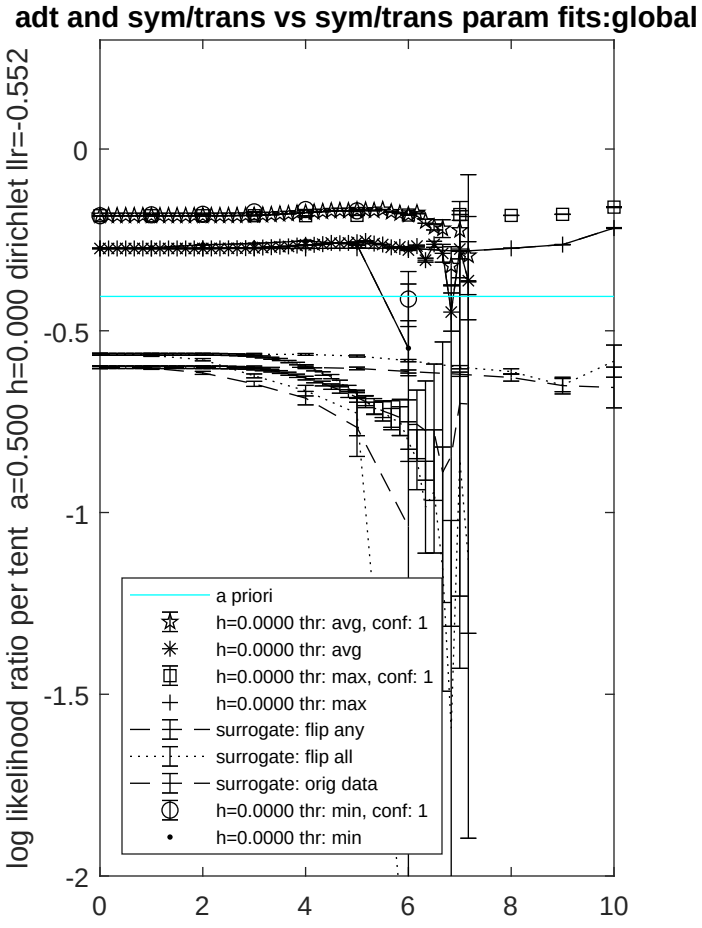

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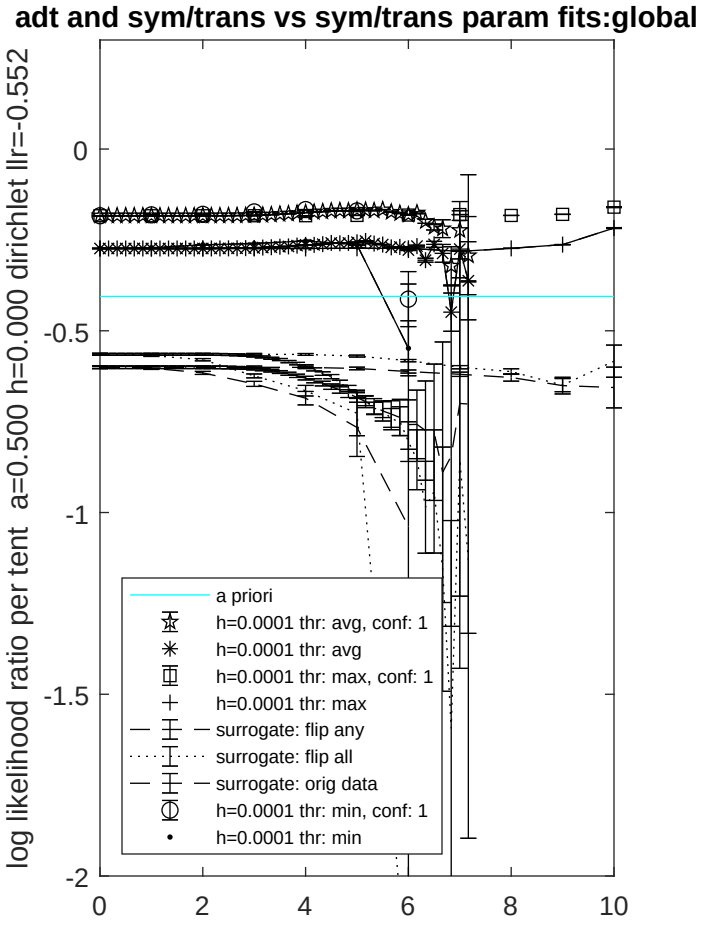

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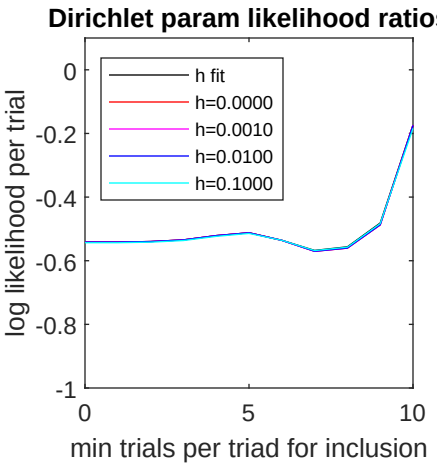

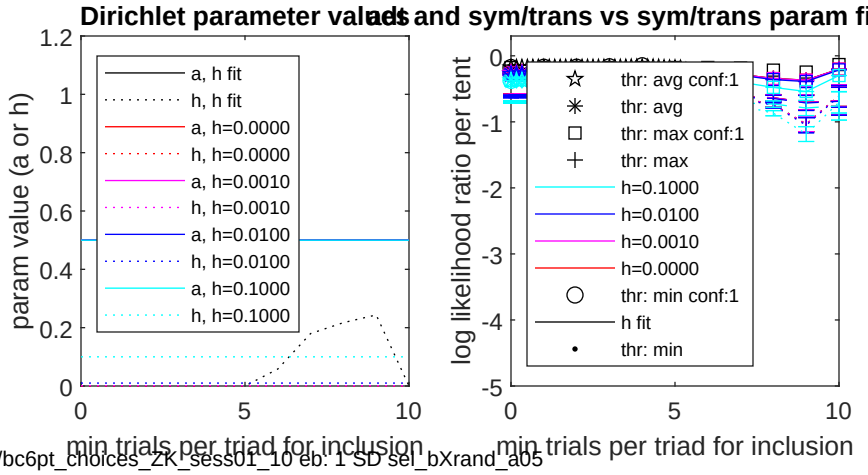

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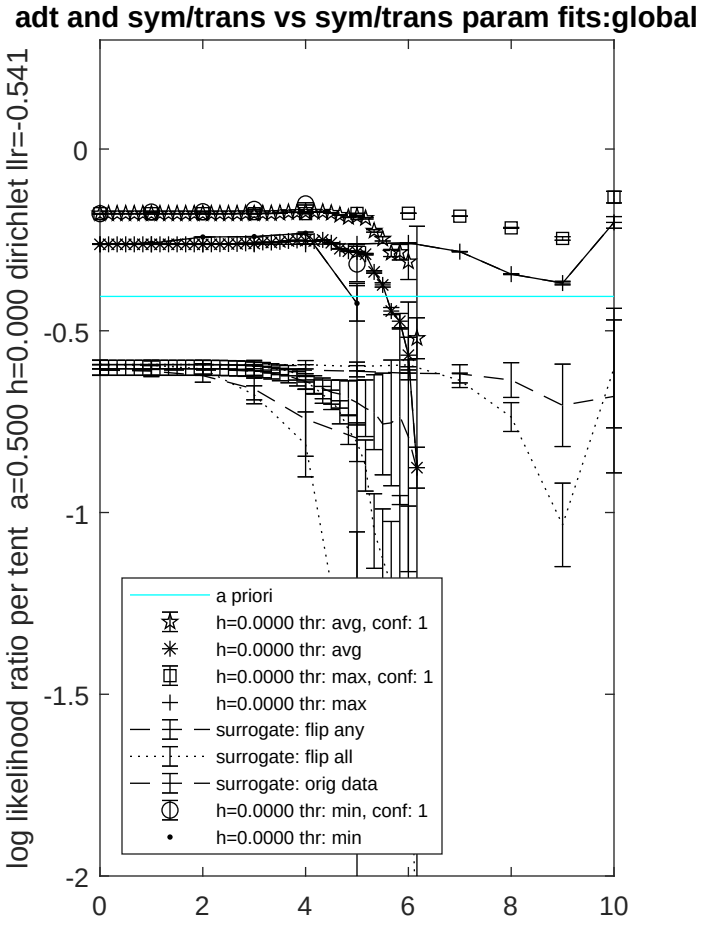

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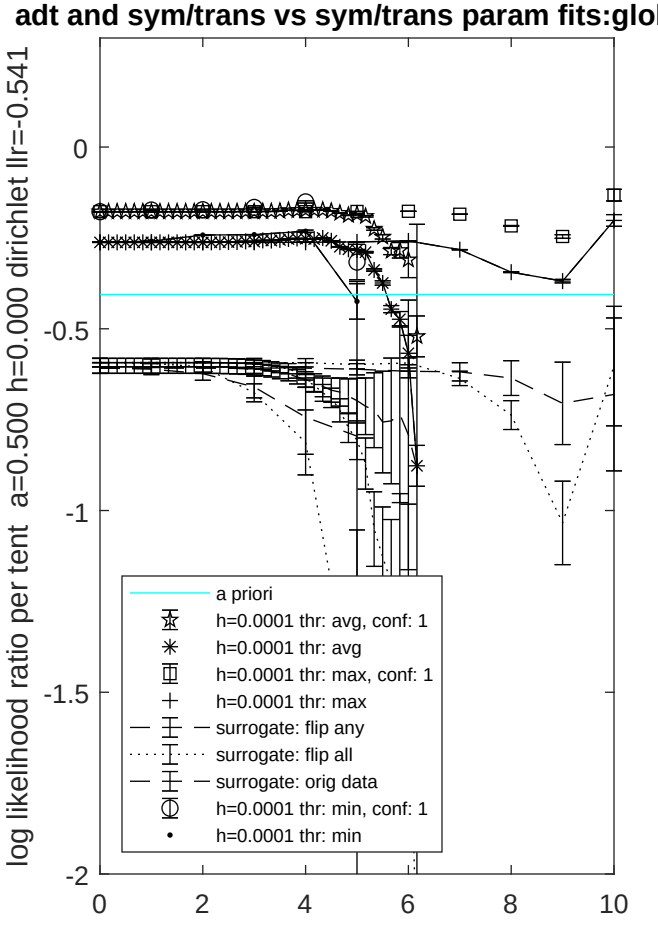

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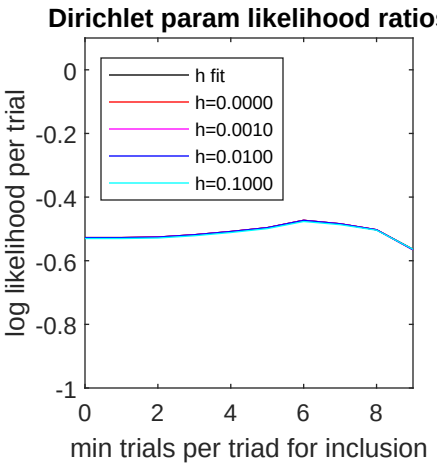

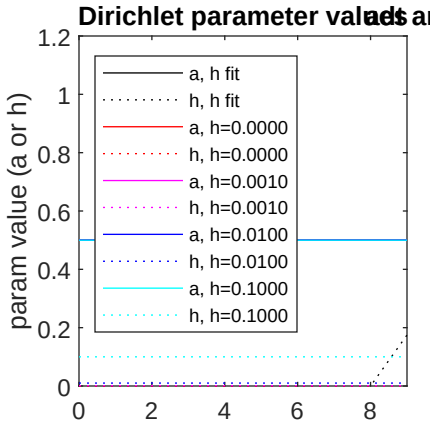

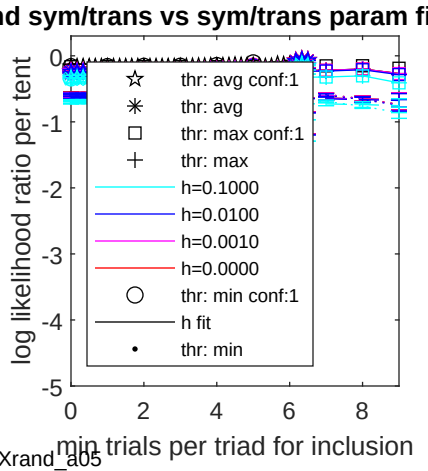

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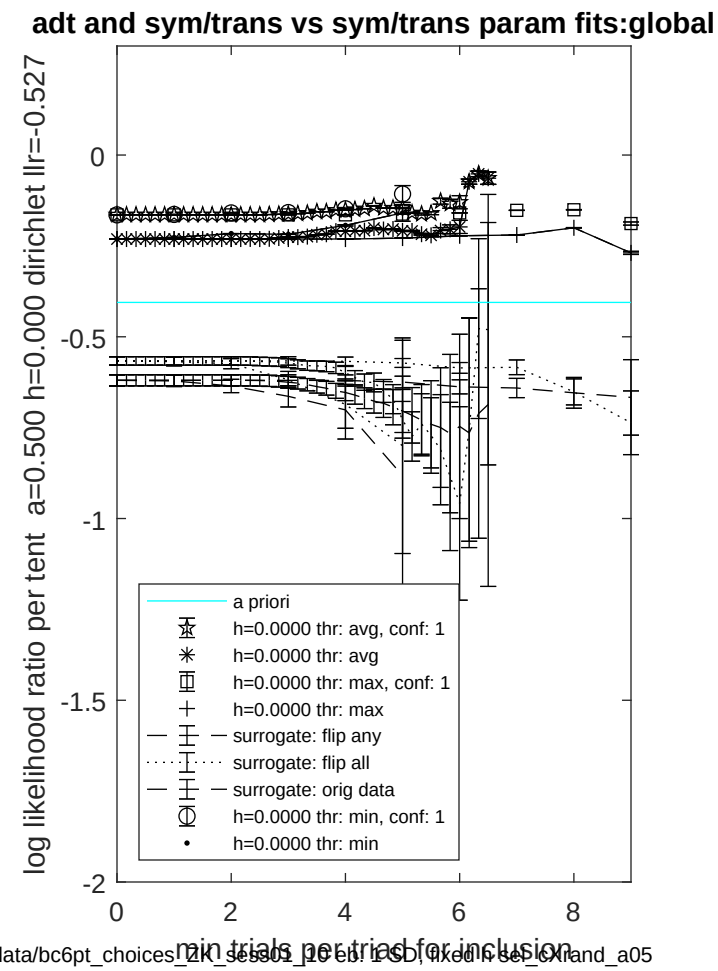

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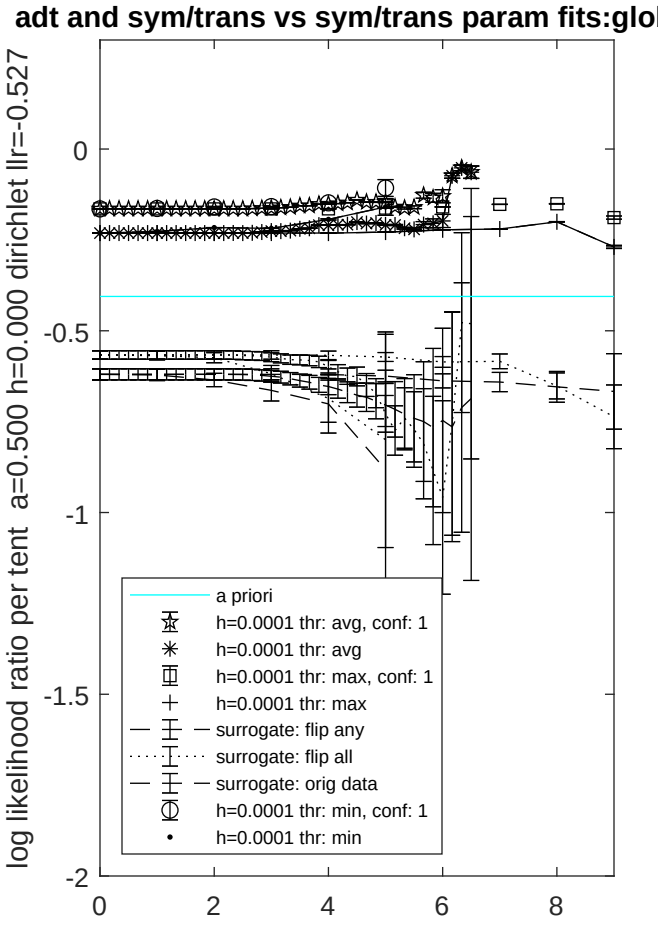

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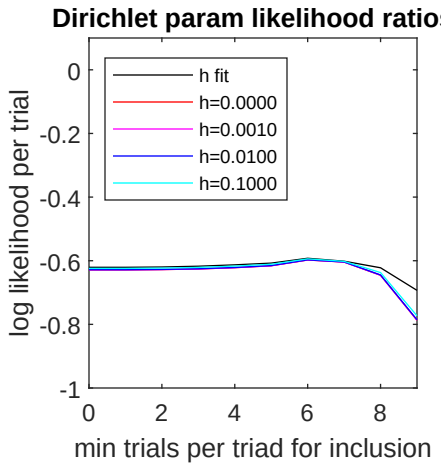

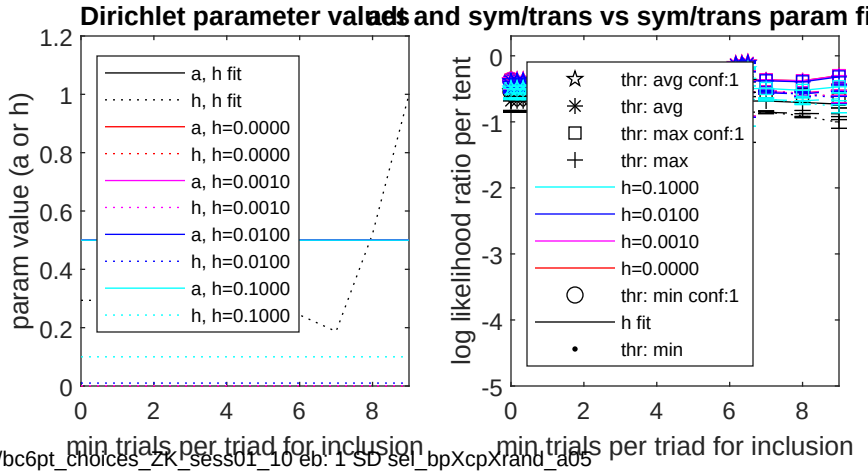

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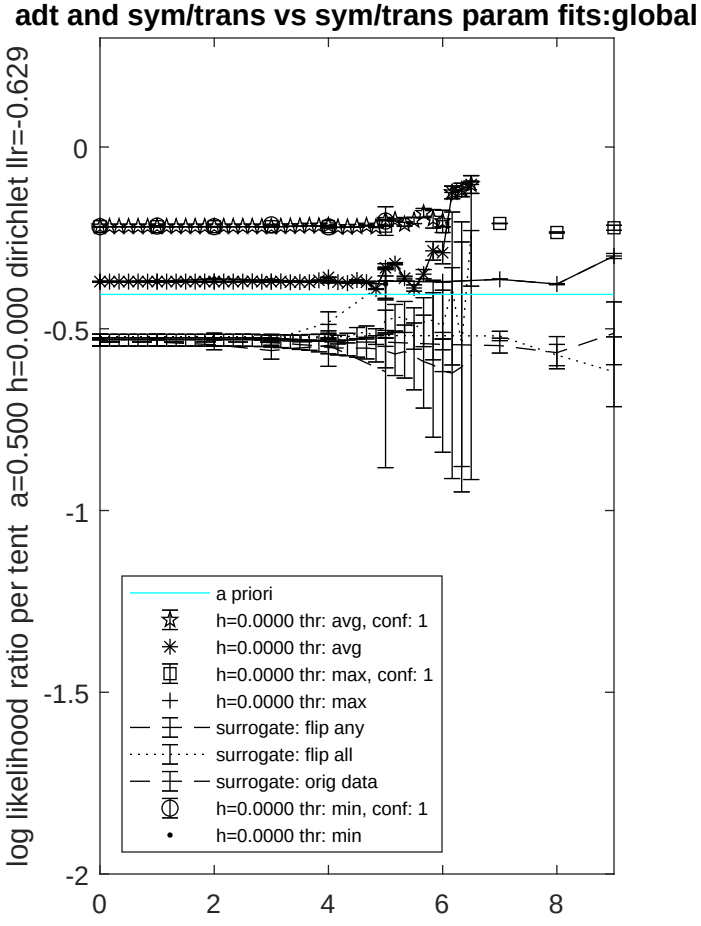

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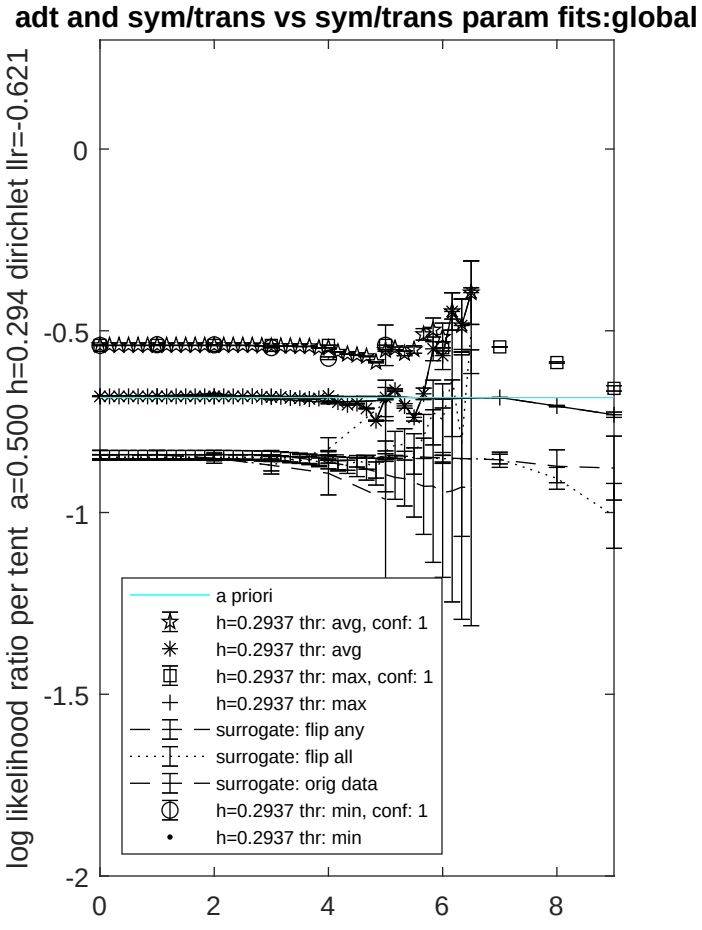

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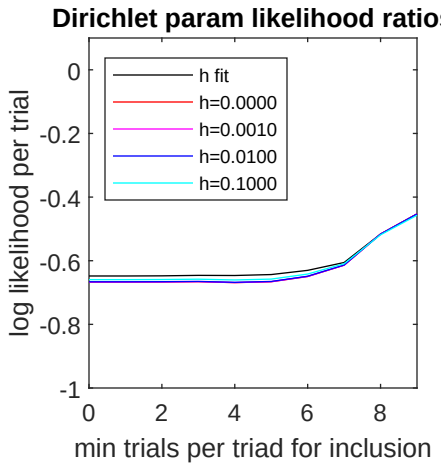

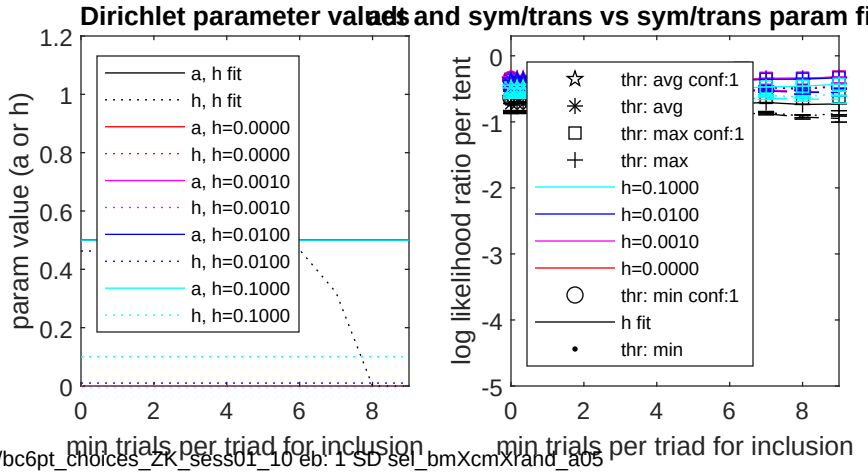

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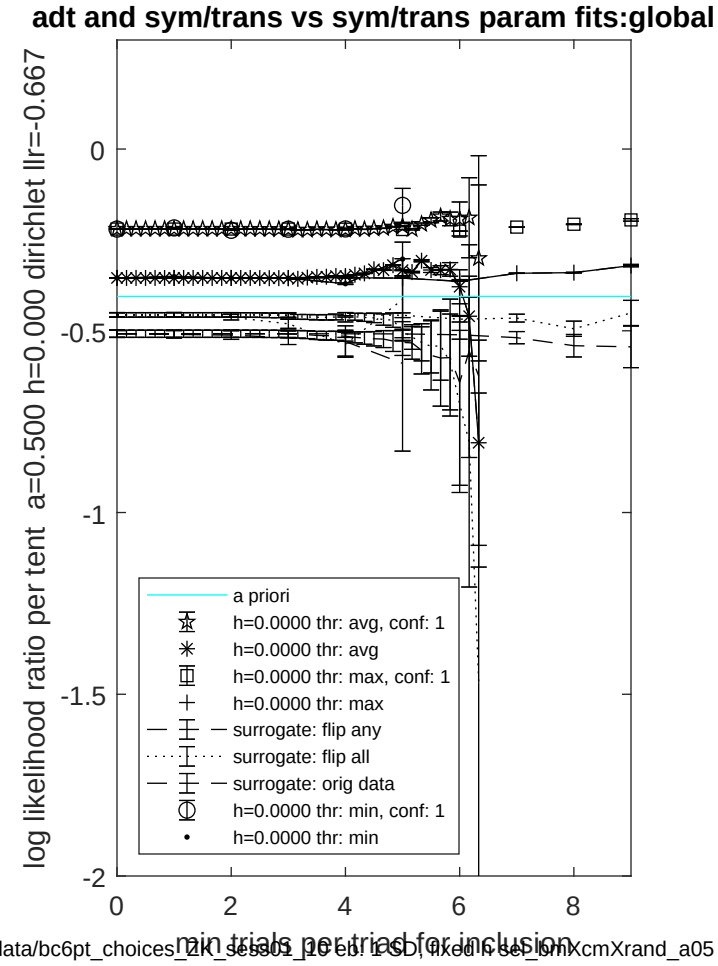
adt and sym/trans vs sym/trans param fits:global
log likelihood ratio per tent a=0.500 h=0.000 dirichlet llr=-0.667
0
-0.5
-1
-1.5
-2
0
2
4
6
8
min trials per triad for inclusion
a priori
h=0.0000 thr: avg, conf: 1
h=0.0000 thr: avg
h=0.0000 thr: max, conf: 1
h=0.0000 thr: max
surrogate: flip any
surrogate: flip all
surrogate: orig data
h=0.0000 thr: min, conf: 1
h=0.0000 thr: min
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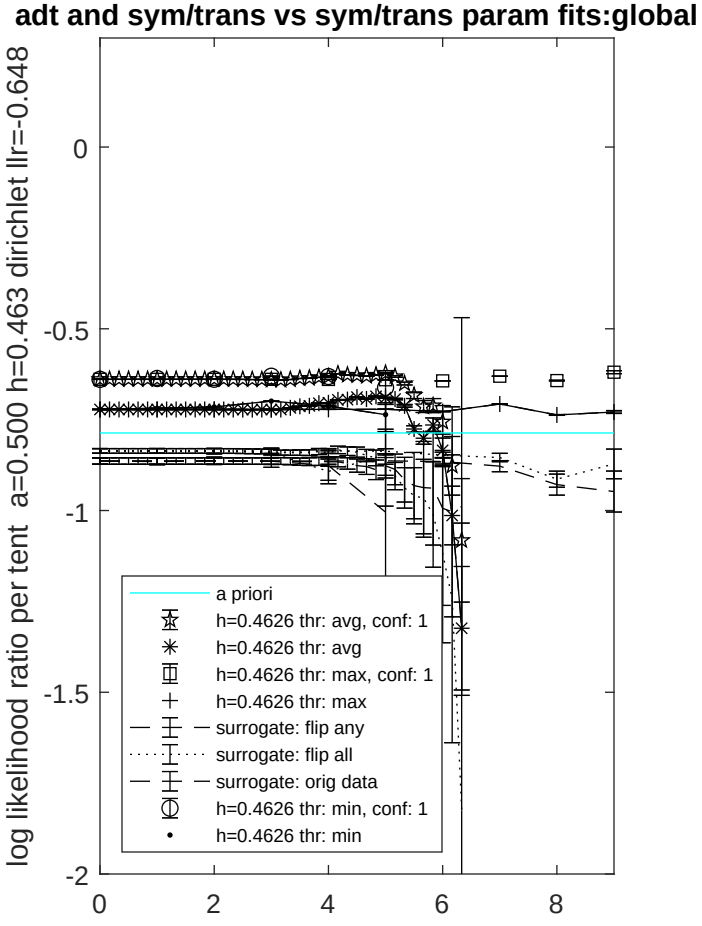

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