

NAV evaluation Harmonization of calving traits

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Innovation

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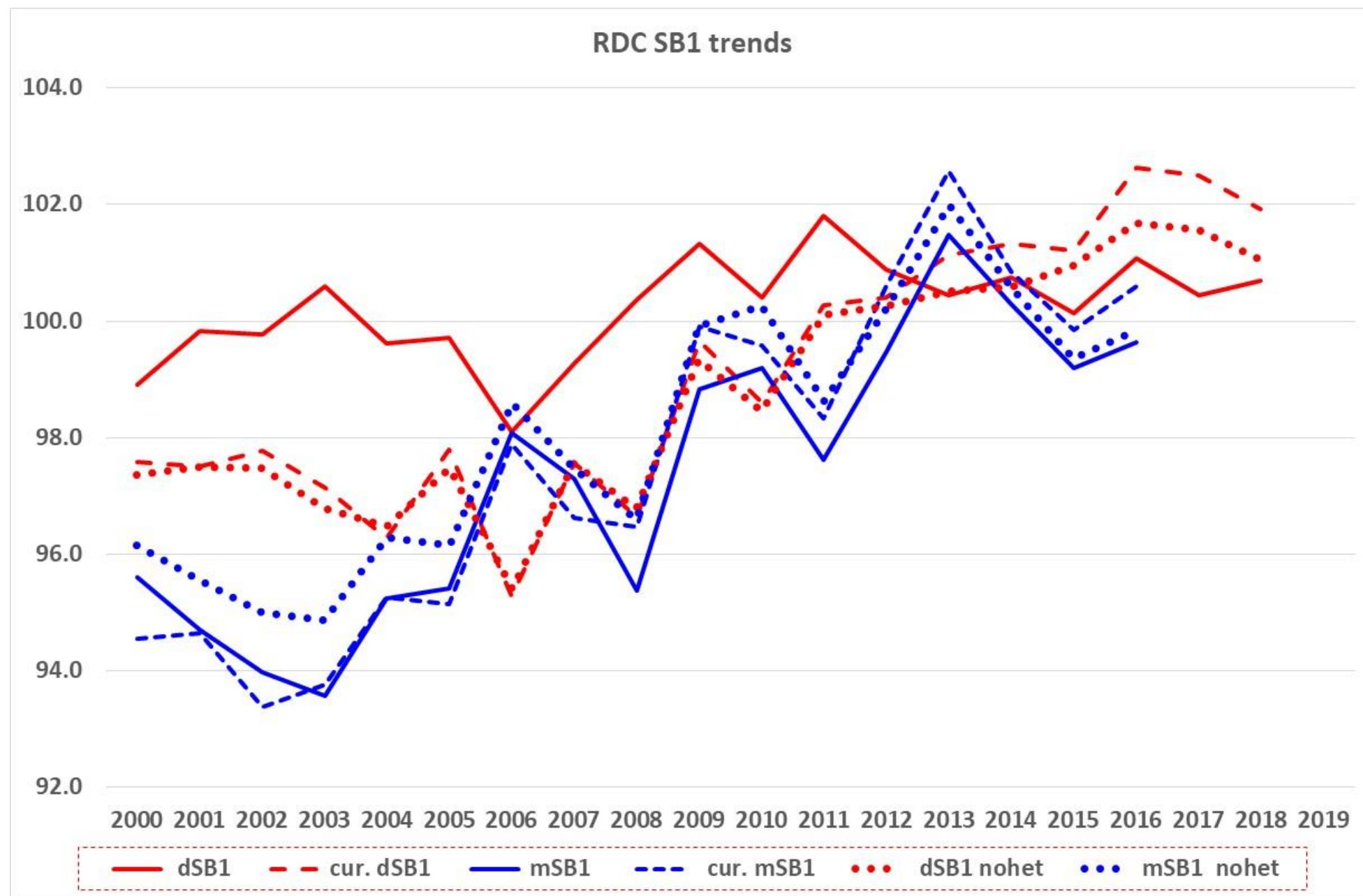
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Mælkeafgiftsfonden

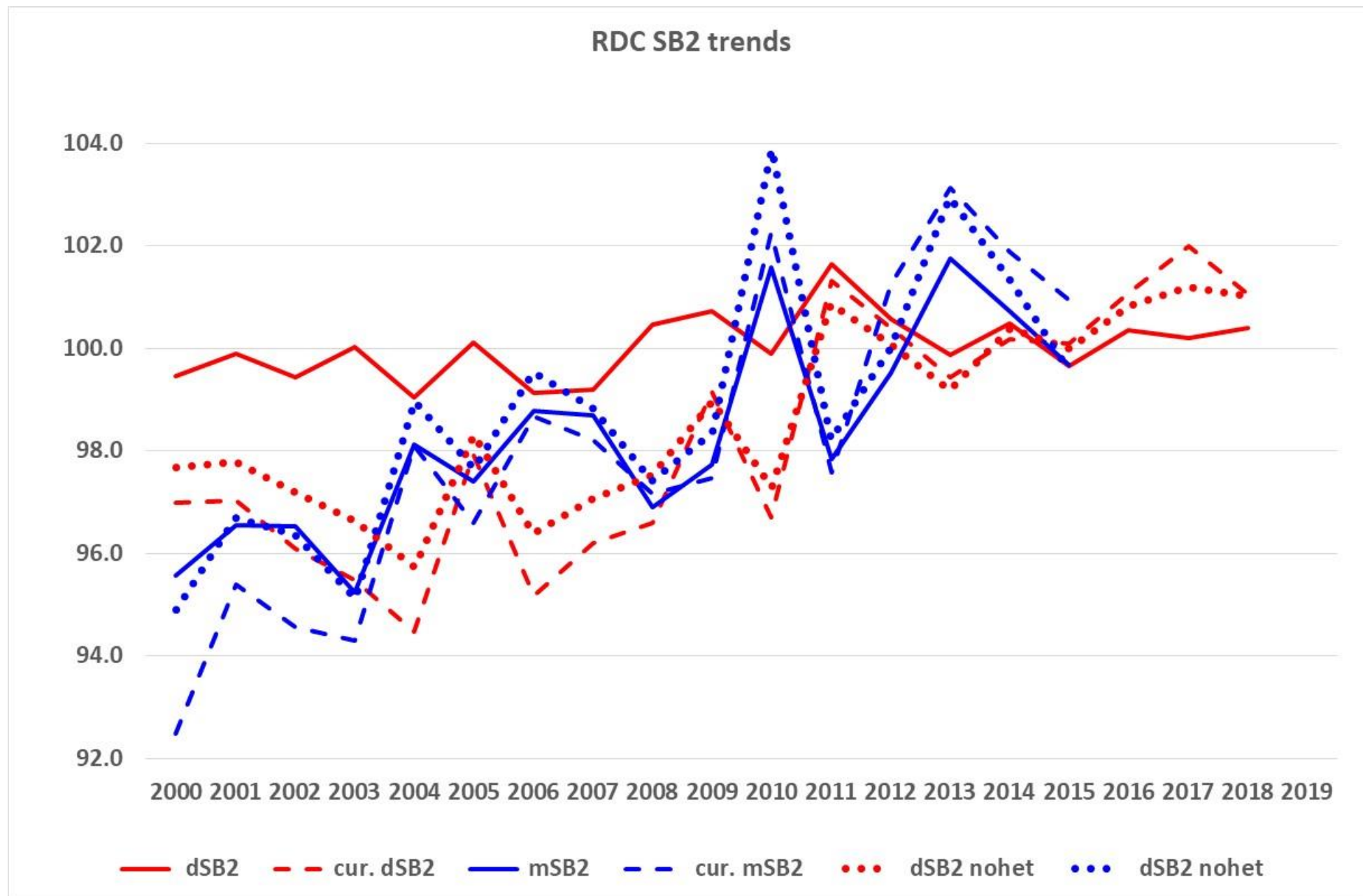
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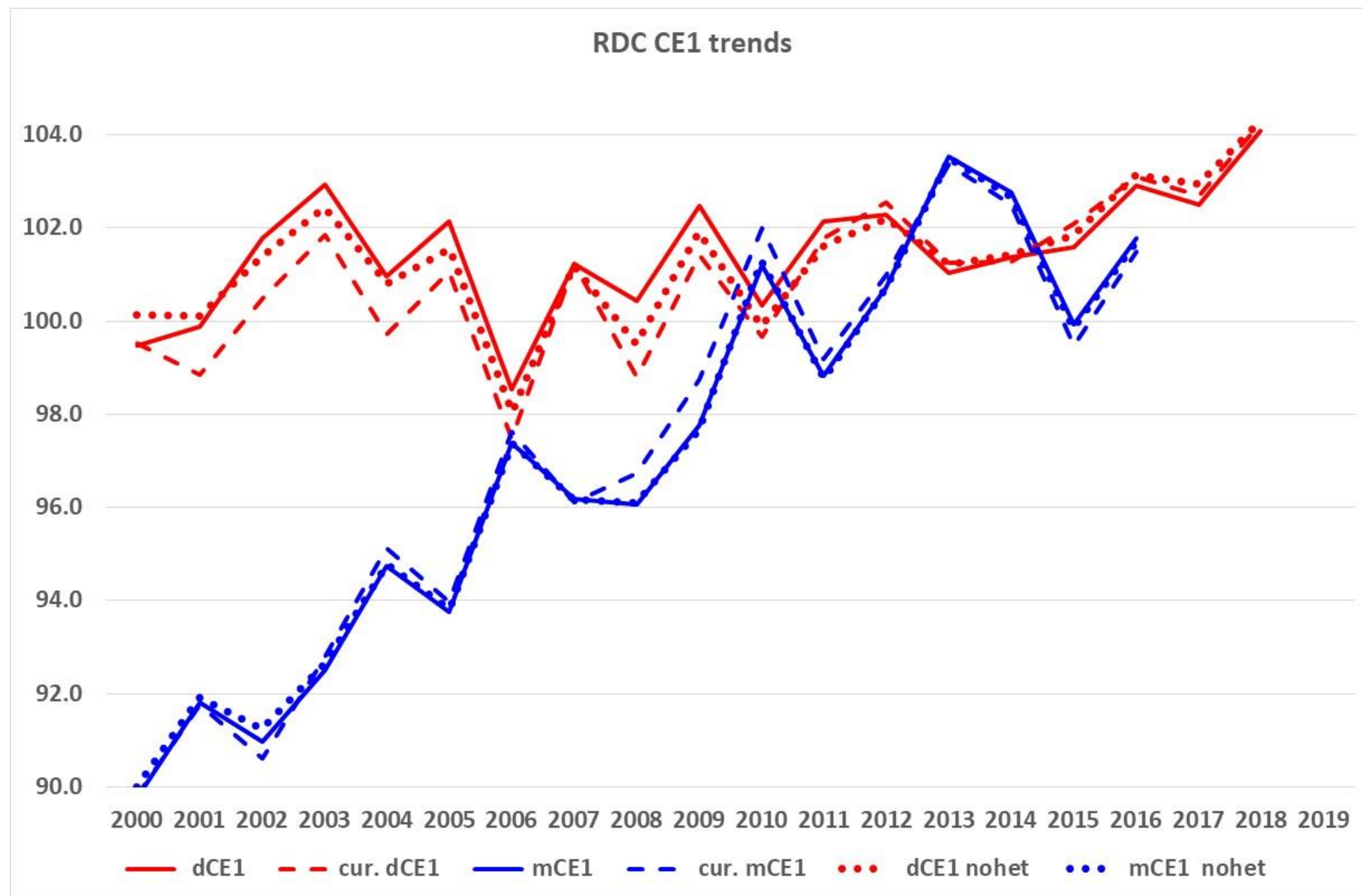
Since last meeting

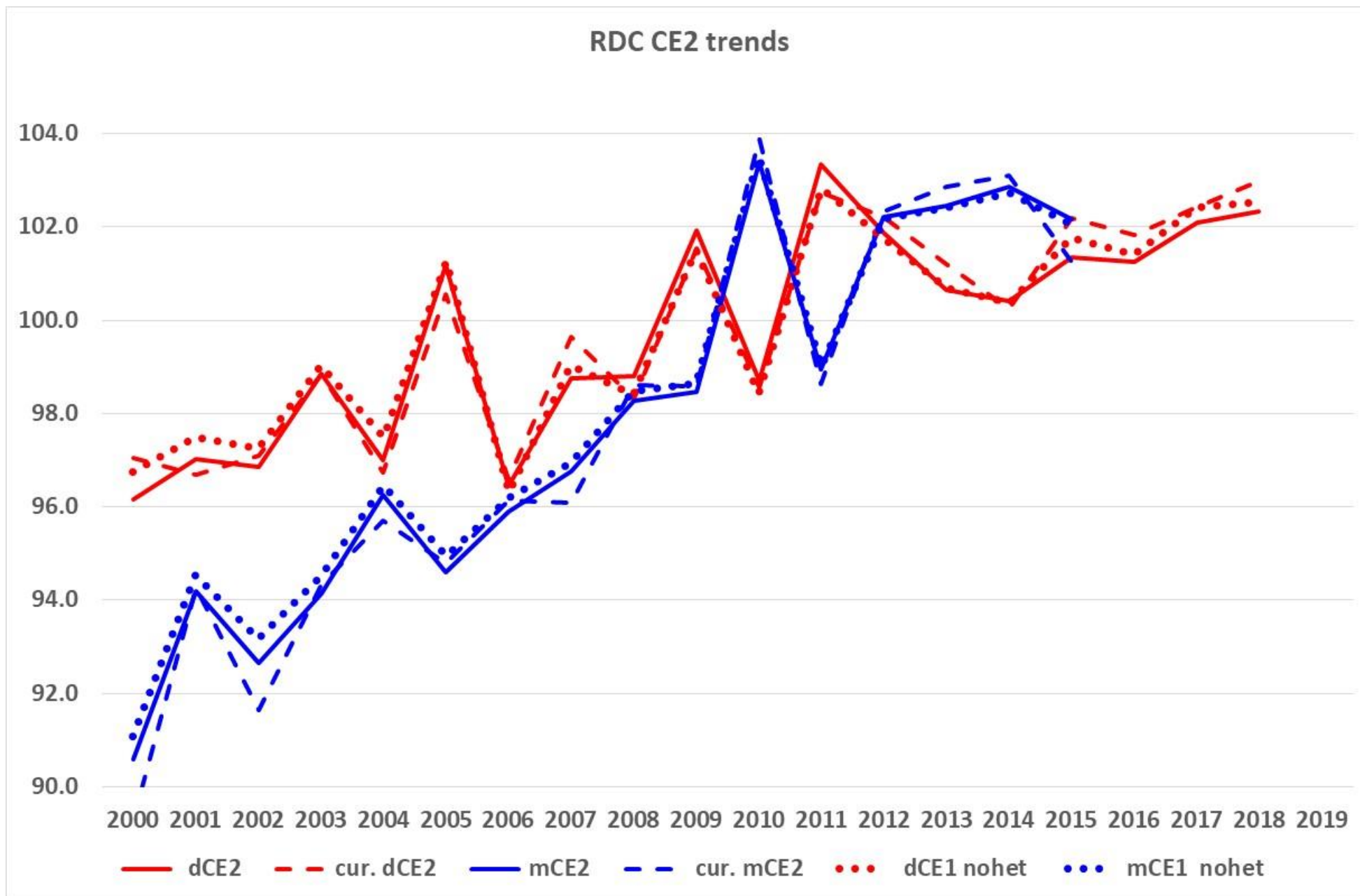
Work on RDC evaluation

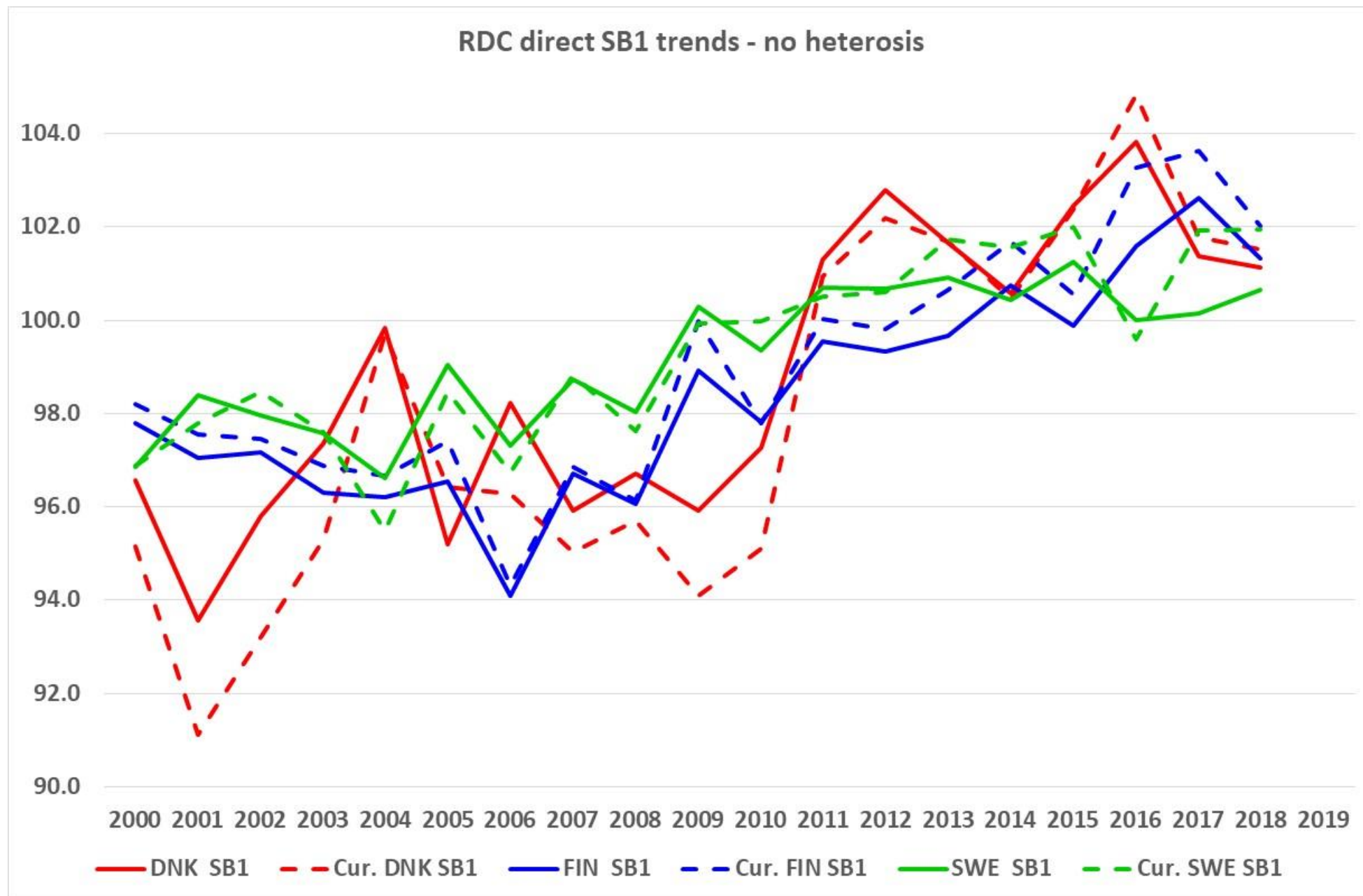
- Sire index trends
- Model with and without heterosis
- Interbull validation
 - JER
 - HOL

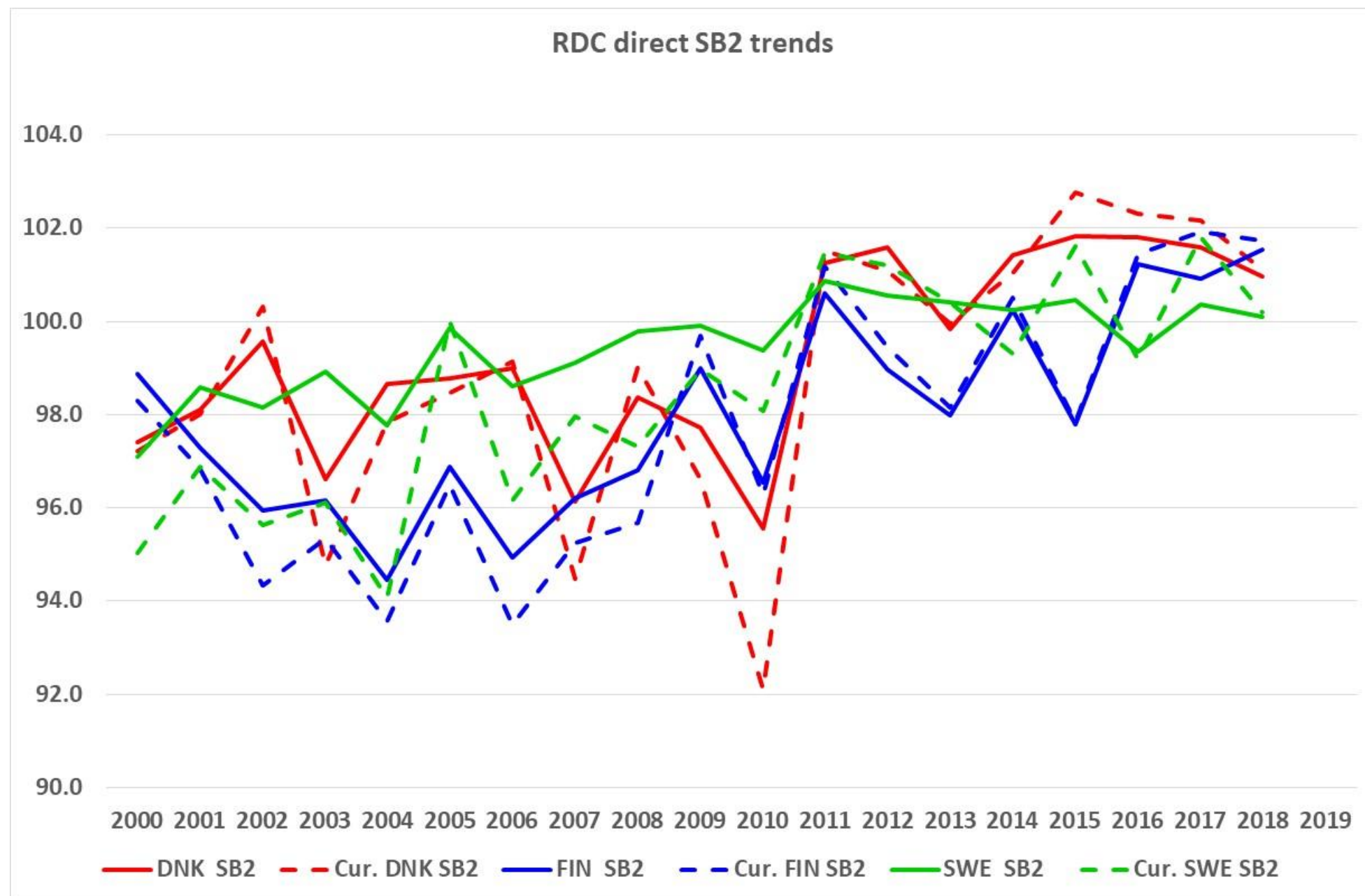












RDC sire-corr. with current, sires born $\geq$ 2000
 $R^2 > 0.7$ for birth idx, > 0.7 for calving index

	With het.	Without heterosis				"Trine" w. het	"Trine" No het.
	All	All	DNK	FIN	SWE	All	All
dSB1	0.84	0.95	0.97	0.96	0.95	0.90	0.98
dCE1	0.96	0.97	0.98	0.98	0.97	0.99	0.99
dSB2	0.87	0.96	0.97	0.97	0.95	0.92	0.97
dCE2	0.92	0.93	0.95	0.95	0.93	0.97	0.97
mSB1	0.92	0.93	0.95	0.93	0.91	0.95	0.95
mCE1	0.97	0.96	0.98	0.98	0.96	0.99	0.99
mSB2	0.82	0.83	0.75	0.85	0.81	0.80	0.81
mCE2	0.95	0.95	0.95	0.97	0.94	0.96	0.96
Birth idx	0.94	0.96	0.99	0.97	0.95		
Calv idx	0.89	0.91	0.93	0.92	0.87		

RDC sire-correlations between model with and without heterosis included

Trines correlation calculations

	Current model		New model	
	Direct	Maternal	Direct	Maternal
SB1	0.9976	0.9962	0.90	1.00
CE1	0.9999	0.9976	1.00	1.00
CS1	0.9998	0.9999	1.00	1.00
SB2	0.9997	0.9982	0.93	1.00
CE2	0.9994	0.9983	1.00	1.00
CS2	0.9999	0.9973	1.00	1.00

RDC heterosis – 1st calving traits

nordhet: srbxfay + srbxnrf + srbxcay + fayxnrf + fayxcay + fayxfic + nrfxcay

resthet: totalhet - nordhet (mostly RDM, ABS, RHF and their crosses with SRB, FAY, NRF and CAY)

Trait	Heterosis effect	New	Current	SD New	SD Current	New index	Current index	Diff. New-current
SB1	nordhet	0.2717	0.0053	0.0876	0.0322	31.0	1.7	29.4
SB1	resthet	0.2345	0.0106	0.0876	0.0322	26.8	3.3	23.5
SB1	mnordhet	-0.0137	0.0073	0.0610	0.0235	-2.2	3.1	-5.4
SB1	mresthet	-0.0017	0.0114	0.0610	0.0235	-0.3	4.9	-5.2
CE1	nordhet	0.1036	-0.0175	0.1536	0.1413	6.7	-1.2	8.0
CE1	resthet	0.0975	-0.0146	0.1536	0.1413	6.3	-1.0	7.4
CE1	mnordhet	0.0309	0.0367	0.1127	0.0939	2.7	3.9	-1.2
CE1	mresthet	0.0407	0.0491	0.1127	0.0939	3.6	5.2	-1.6

RDC heterosis – Later calving traits

nordhet: srbxfay + srbxnrf + srbxcay + fayxnrf + fayxcay + fayxfic + nrfxcay

resthet: totalhet - nordhet (mostly RDM, ABS, RHF and their crosses with SRB, FAY, NRF and CAY)

Trait	Heterosis effect	New	Current	SD New	SD Current	New index	Current index	Diff. New-current
SB2	nordhet	0.1206	0.0036	0.0450	0.0145	26.8	2.5	24.3
SB2	resthet	0.1044	0.0028	0.0450	0.0145	23.2	1.9	21.3
SB2	mnordhet	-0.0013	0.0028	0.0302	0.0134	-0.4	2.1	-2.5
SB2	mresthet	-0.0010	0.0044	0.0302	0.0134	-0.3	3.3	-3.6
CE2	nordhet	0.0489	0.0058	0.0810	0.0660	6.0	0.9	5.2
CE2	resthet	0.0402	-0.0071	0.0810	0.0660	5.0	-1.1	6.0
CE2	mnordhet	0.0239	0.0201	0.0696	0.0542	3.4	3.7	-0.3
CE2	mresthet	0.0179	0.0228	0.0696	0.0542	2.6	4.2	-1.6

Summary

- Very high (too high) effect of heterosis for direct SB1 and SB2.
- In new: Correlations between with and without heterosis included are low for direct SB1 and SB2 – otherwise close to 1.00
- In current: Correlations between with and without heterosis included are close to 1.00
- Heterosis could be omitted !
- BUT: There are still problems maternal SB2