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**Supplementary file S5:** Variance-covariance matrix of the fixed effects of the developed multi-response biomass model for Norway spruce.

Effect		Stem			Branches		Foliage		Stump		Roots		
		<i>Incept</i>	$d_s/(d_s+k)$	$\ln(h)$	<i>Incept</i>	$\ln(d_s)$	<i>Incept</i>	$d_s/(d_s+k)$	<i>Incept</i>	$d_s/(d_s+k)$	<i>Incept</i>	$d_s/(d_s+k)$	$\ln(h)$
Stem	<i>Incept</i>	<b>0.1641</b>	<b>-0.3517</b>	<b>0.07347</b>	-0.00718	0.008252	0.002021	-0.00332	0.0185	-0.02768	-0.00985	0.01876	-0.01005
	$d_s/(d_s+k)$	<b>-0.3517</b>	<b>0.7646</b>	<b>-0.1667</b>	0.01255	-0.0156	-0.00386	0.006482	-0.03324	0.05148	0.02867	-0.05077	0.02281
	$\ln(h)$	<b>0.07347</b>	<b>-0.1667</b>	<b>0.04674</b>	6.4E-05	-0.00008	0.000192	-0.00032	-0.00168	0.001725	-0.016	0.02593	-0.00641
Branches	<i>Incept</i>	-0.00718	0.01255	6.4E-05	<b>0.00838</b>	<b>-0.00854</b>	0.01008	-0.01581	0.01693	-0.02348	0.01333	-0.01848	0.00003
	$\ln(d_s)$	0.008252	-0.0156	-0.00008	<b>-0.00854</b>	<b>0.01061</b>	-0.01171	0.01966	-0.01993	0.0292	-0.01569	0.02298	-0.00004
Foliage	<i>Incept</i>	0.002021	-0.00386	0.00019	0.01008	-0.01171	<b>0.03777</b>	<b>-0.062</b>	0.02607	-0.03755	0.04401	-0.063	-0.00056
	$d_s/(d_s+k)$	-0.00332	0.00648	-0.00032	-0.01581	0.01966	<b>-0.062</b>	<b>0.1041</b>	-0.04298	0.06303	-0.07252	0.1058	0.000947
Stump	<i>Incept</i>	0.0185	-0.03324	-0.00168	0.01693	-0.01993	0.02607	-0.04298	<b>0.2289</b>	<b>-0.3212</b>	0.1513	-0.2125	0.000568
	$d_s/(d_s+k)$	-0.02768	0.05148	0.00173	-0.02348	0.0292	-0.03755	0.06303	<b>-0.3212</b>	<b>0.4586</b>	-0.2121	0.3033	-0.00073
Roots	<i>Incept</i>	-0.00985	0.02867	-0.016	0.01333	-0.01569	0.04401	-0.07252	0.1513	-0.2121	<b>0.6013</b>	<b>-0.941</b>	<b>0.1671</b>
	$d_s/(d_s+k)$	0.01876	-0.05077	0.02593	-0.01848	0.02298	-0.063	0.1058	-0.2125	0.3033	<b>-0.941</b>	<b>1.4877</b>	<b>-0.2751</b>
	$\ln(h)$	-0.01005	0.02281	-0.00641	0.00003	-0.00004	-0.00056	0.000947	0.00057	-0.00073	<b>0.1671</b>	<b>-0.2751</b>	<b>0.06951</b>

Off diagonal blocks consist of the across equation covariances of the fixed effects, and block diagonal is variances-covariances matrix of the sub-model for the individual tree component (bold).