

Table S2. Mean molecular distances between groups. Standard errors are in grey.

<i>zmp/zmp</i>	<i>iga</i>	<i>zmpC</i>	<i>zmpB</i>	<i>zmpD</i>
<i>iga</i>		0.018	0.025	0.029
<i>zmpC</i>	0.708		0.025	0.029
<i>zmpB</i>	0.951	0.951		0.014
<i>zmpD</i>	0.943	0.959	0.566	