

Table S3. Mean molecular distances between ZmpB subgroups and the ZmpD cluster. Standard errors are in grey.

	<i>zmpB_1</i>	<i>zmpB_2</i>	<i>zmpB_3</i>	<i>zmpB_4</i>	<i>zmpB_5</i>	<i>zmpB_6</i>	<i>zmpB_7</i>	<i>zmpB_8</i>	<i>zmpB_9</i>	<i>zmpB_10</i>	<i>zmpD_11</i>
<i>zmpB_1</i>		0.010	0.009	0.011	0.015	0.014	0.015	0.017	0.018	0.023	0.013
<i>zmpB_2</i>	0.266		0.005	0.012	0.015	0.014	0.013	0.018	0.018	0.019	0.014
<i>zmpB_3</i>	0.225	0.089		0.009	0.016	0.014	0.013	0.019	0.018	0.021	0.014
<i>zmpB_4</i>	0.301	0.353	0.231		0.013	0.014	0.016	0.017	0.020	0.025	0.011
<i>zmpB_5</i>	0.504	0.480	0.508	0.422		0.013	0.015	0.022	0.021	0.027	0.012
<i>zmpB_6</i>	0.440	0.446	0.444	0.439	0.442		0.016	0.023	0.020	0.024	0.013
<i>zmpB_7</i>	0.417	0.416	0.419	0.461	0.521	0.486		0.020	0.018	0.024	0.015
<i>zmpB_8</i>	0.629	0.603	0.612	0.634	0.674	0.679	0.577		0.014	0.022	0.018
<i>zmpB_9</i>	0.619	0.617	0.625	0.627	0.681	0.653	0.574	0.411		0.021	0.017
<i>zmpB_10</i>	0.762	0.726	0.736	0.751	0.811	0.772	0.739	0.681	0.687		0.023
<i>zmpD_11</i>	0.399	0.459	0.458	0.378	0.463	0.461	0.502	0.660	0.660	0.810	