

**Table S4.** Top 20 differentially methylated genes among pancreatic cancer subtypes

<b>Gene Name</b>	<b>DE-<math>\beta</math></b>	<b>P</b>	<b>adjustP</b>
<i>MIR135B</i>	-0.2454	5.15E-28	4.02E-26
<i>CAPN8</i>	-0.2683	4.45E-25	1.16E-23
<i>FAM25A</i>	-0.1329	4.27E-25	1.16E-23
<i>CAPN2</i>	-0.1055	3.46E-24	5.39E-23
<i>HRH1</i>	-0.1649	1.10E-22	1.44E-21
<i>MTMR7</i>	0.1595	1.72E-21	1.91E-20
<i>SLAMF7</i>	-0.1753	4.27E-20	4.17E-19
<i>GSDMC</i>	-0.2042	4.85E-20	4.21E-19
<i>S100A10</i>	-0.1218	8.34E-20	6.51E-19
<i>ARNTL2</i>	-0.1006	3.62E-19	2.57E-18
<i>CCDC70</i>	-0.1410	1.63E-16	9.79E-16
<i>SNORD114.29</i>	-0.1990	1.63E-16	9.79E-16
<i>SLPI</i>	-0.1375	1.90E-16	1.06E-15
<i>MYEOV</i>	-0.1251	5.30E-16	2.76E-15
<i>HOXC12</i>	0.1022	1.18E-15	5.13E-15
<i>UCA1</i>	-0.1927	1.18E-15	5.13E-15
<i>KRTAP2.4</i>	-0.2117	1.85E-15	7.61E-15
<i>LOC728392</i>	0.2163	3.42E-15	1.34E-14
<i>SHROOM3</i>	-0.1247	5.67E-15	2.11E-14
<i>ISG20L2</i>	-0.1166	6.08E-15	2.16E-14