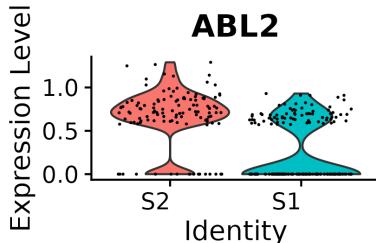
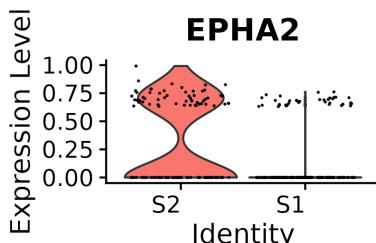
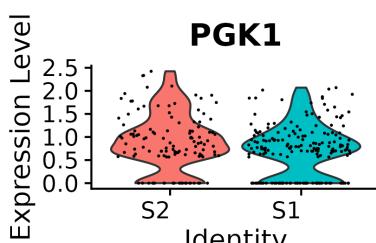
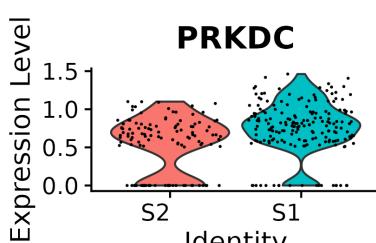
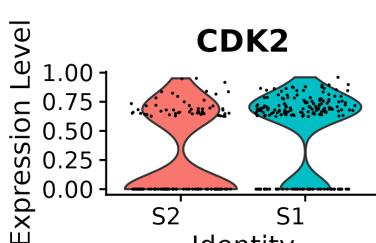
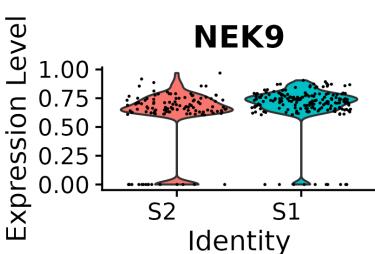


Log Fold change in genes between Sample 1 cells vs Sample 2 cells

Gene_Name	avg_log2FC	pct.1	pct.2	violinPlot
ABL2	0.5543281	0.857	0.398	 <p>ABL2</p> <p>Expression Level</p> <p>S2 S1</p> <p>Identity</p>
EPHA2	0.3163869	0.462	0.199	 <p>EPHA2</p> <p>Expression Level</p> <p>S2 S1</p> <p>Identity</p>
PGK1	0.3131115	0.815	0.715	 <p>PGK1</p> <p>Expression Level</p> <p>S2 S1</p> <p>Identity</p>
PRKDC	-0.3067348	0.731	0.898	 <p>PRKDC</p> <p>Expression Level</p> <p>S2 S1</p> <p>Identity</p>
CDK2	-0.2900257	0.445	0.737	 <p>CDK2</p> <p>Expression Level</p> <p>S2 S1</p> <p>Identity</p>

Gene_Name	avg_log2FC	pct.1	pct.2	violinPlot
WNK4	-0.2475849	0.311	0.538	<p>WNK4</p>
EGFR	-0.2316553	0.622	0.823	<p>EGFR</p>
AXL	0.2289109	0.866	0.634	<p>AXL</p>
EPHB2	-0.1929875	0.345	0.522	<p>EPHB2</p>
RIPK2	0.1825412	0.681	0.538	<p>RIPK2</p>

Gene_Name	avg_log2FC	pct.1	pct.2	violinPlot
STK25	-0.1351858	0.916	0.978	<p>STK25</p>
DTYMK	-0.1309976	0.630	0.704	<p>DTYMK</p>
LYN	0.1241359	0.429	0.317	<p>LYN</p>
RPS6KA3	-0.1157113	0.950	0.984	<p>RPS6KA3</p>
DDR1	-0.1055468	0.773	0.892	<p>DDR1</p>

Gene_Name	avg_log2FC	pct.1	pct.2	violinPlot
NEK9	-0.1046200	0.866	0.946	
LZTR1	0.1045338	0.958	0.898	