Supplemental Table S2. Multiple Reaction Montitoring Peptides and Transitions			
Protein	**parent ion m/z**	**Q3 fragment m/z**	**sequence**
CYCLIN K1/K2	674.4	644.8	30 DLAHTPSQLEGLDPATEAR 48
		759.7	
		929.8	
	477.9	694.4	56 FIFDVGTR 63
		432.6	
		547.4	
CYCLIN K1 unique	785.8	604.2	375 KPPLAAALGEAEPPGPVDATDLPK 398
		1148.9	
		1206.8	
	625.0	581.8	358 IETTHPPLPPAHPPPDR 374
		646.2	
		889.8	
CYCLIN K2 unique	709.7	744.4	319 QPATHLLPSPLEDSLLCPR 337
		860.6	
		910.9	
		928.7	
		1102.8	
CYCLIN L1	698.0	770.9	348 AEEKSPISINVK 359
		839.8	
		967.9	
CYCLIN T1	476.2	504.8	27 FGVDPDKELSYR 38
		667.8	
		640.4	
	910.8	952.0	351 TSENLALTGVDHSLPQDGSNAFISQK 376
		1291.9	
		1058.3	
	630.5	702.4	439 MPIEGSENPERPFLEK 454
		766.9	
		871.7	
CYCLIN T2	674.9	1008.6	80 NIISSTALFLAAK 92
		1121.9	
		1064.6	
CDK9	512.9	471.6	295 LLVLDPAQR 303
		586.4	
		798.6	
	581.9	716.6	285 DPYALDLIDK 294
		787.8	
		950.9	
	591.9	884.5	25 IGQGTFGEVFK 35
		1069.7	
		827.4	
CDK11	680.5	1060.8	648 IWPGYSELPAVK 659
		963.8	
		531.1	
	510.4	686.5	462 EGFPITSLR 472
		589.6	
		476.5	
	643.6	1043.6	431 IEEGTYGVVYR 441
		914.8	
		756.7	
CDK12	676.9	783.8	1052 QSGVVVEEPPPSK 1064
		882.7	
		981.8	
	455.8	494.6	611 TQVSVTAAIPHLK 623
		607.8	
		678.8	
	794.4	855.8	381 HSSISPVRLPLNSSLGAELSR 401
		981.5	
		889.9	
CDC2L5 (CDK13)	734.9	1153.9	1056 TNTPQGVLPSSQLK 1069
		772.6	
		659.3	
	795.8	1121.1	1110 VETDAAQAAVQ SAFAVLLTQL IK 1132
		1064.6	
		837	
	628.7	759.7	1240 AALLQLLAQHQPQDDPK 1256
		815.9	
		695.4	

Supplemental Table S2. Multiple Reaction Monitoring (MRM) peptides and transitions used for the detection of CDK and CYCLIN proteins.