	SAM HG4: 1-3MBT-31-3MBT-4/SAMD1/SAMD7 SAM-1 SAM HG4: 1-3MBT-31-3MBT-4/SAMD1/SAMD7 METO SAM HG4: 1-3MBT-31-3MBT-4/SAMD1/SAMD7 METO SAM HG4: 1-3MBT-31-3MBT-4/SAMD1/SAMD7 ZHCZHC SAM HG4: 1-3MBT-31-3MBT-4/SAMD1/SAMD7 ZHCZHC SAM HG4: 1-3MBT-31-3MBT-4/SAMD1/SAMD7 SAM-2 SAM HG4: 1-3MBT-31-3MBT-4/SAMD1/SAMD7 SAM-2 SAM HG4: 1-3MBT-31-3MBT-4/SAMD1/SAMD7 RFM 1 SAM HG4: 1-3MBT-31-3MBT-4/SAMD1/SAMD7 RFM 1 SAM HG4: 1-3MBT-31-3MBT-4/SAMD1/SAMD7 RFM 1 SAM HG4: 1-3MBT-31-3MBT-4/SAMD1/SAMD7 RFB SAMD1/SAMD7 RFB SAMD1/SAMD1/SAMD7 RFB SAMD1/SAMD1/SAMD7 RFB SAMD1/SAMD1/SAMD7 RFB SAMD1/SAMD7 RFB SAMD1/SAMD1/SAMD7 RFB SAMD1/SAMD1/SAMD7 RFB
	SAM HG5.8: EPHA1/EPHA2/EPHA3/EPHA4/EPH. Gene presence SAM HG5.8: EPHA1/EPHA2/EPHA3/EPHA4/EPH. SAM HG5.8: EPHA1/EPHA2/EPHA3/EPHA4/EPH. Pkinase Tyr SAM HG5.8: EPHA1/EPHA2/EPHA3/EPHA4/EPH. SAM A_TM SAM HG5.8: EPHA1/EPHA2/EPHA3/EPHA4/EPH. EPANA_TM SAM HG5.8: EPHA1/EPHA2/EPHA3/EPHA4/EPH. BAM FG0, IIII IIII IIII IIII IIII IIII IIII I
	SAM I G3 14 III I I I I I I I I I I I I I I I I
•	SAW-10; 3: SAW-10; SAW-10; SAM-10; SAM-10; 3: SAW-10; SAM-10; 3: SAW-10; SAW-10; SAW-10; 3: SAW-10;
	SAM HGS. 1; like: EPHA/EPHAZ/EPHAX/EPHA. Gene presence SAM HGS. 1; like: EPHA/EPHAZ/EPHAX/EPHA. SAM   SAM HGS. 1; like: EPHA/EPHAZ/EPHAX/EPHA. SAM   SAM HGS. 1; like: EPHA/EPHAZ/EPHAX/EPHAX SAM HGS. 1; like: EPHA/EPHAZ/EPHAX/EPHAX/EPHAX SAM HGS. 1; like: EPHA/EPHAZ/EPHAX/EPHAX/EPHAX
	\$AM.HG11.1LR\$AM1 LRR \$ \$AM.HG11.1LR\$AM1 SAM1 SAM1 SAM1 SAM1 SAM1 SAM1 SAM1 S
	 SAM HG3.1 IIKe: ANKS1 A/ANKS1 B/CASKIN1/   Gene presence SAM HG3.1 IIKE: ANKS1 AANKS1 B/CASKIN1/   SAM T2 SAM HG3.1 IIKE: ANKS1 AANKS1 B/CASKIN1/   SAM T2 SAM HG3.1 IIKE: ANKS1 AANKS1 B/CASKIN1/   Interfase H2C2