SKCM TP BRAF WT Correlation: 0.04058904 p-value: 0.819713 1.4 -0 0 0 0 0 1.2 -0 0 0 Metastatic score 0 0 1.0 -0 0 0.8 -0.6 2 4 6 8 10 12 14 16 18 20 22 24 26 28 30 32 34 36 38 40 42 44 46 48 50 52 54 56 58 60 62 64 66 68 70 72 74 76 78

Total mutations contributed by S1