rithess distributions for Representative furnors 1.0 -0.8 0.6 mean\_fitness 0.4 0.2 0.0 NMI\_MGH\_0 -- 9\_HDM\_IMN MI\_MGH\_0 -MI\_MGH\_3 -Met\_MGH\_0 -Met\_MGH\_3 NMI\_MGH\_5 MI\_MGH\_1 MI\_MGH\_2 Met\_MGH\_1 Met\_MGH\_2 NMI\_MGH\_2 NMI\_MGH\_3 NMI\_MGH\_4 tumor