CIFAR100 ResNet18 w=32 1.0 Before permutation Before permutation REPAIR 0.8 -Weight matching (top-1) Activation matching Sinkhorn weight matching Loss Sinkhorn direct 0.4 -0.2 -

0.2

0.6

Interpolation α

0.8

1.0

0.0

0.0

0.0

0.2

0.4

0.6

Interpolation α

0.8

1.0