

Suppl. Table III siRNA duplexes

AKT1	CATGAACGAGTTTGAGTACCT CACATTAAGATCACAGACTTC GATCATGCAGCATCGCTTCTT CAAGATGACAGCATGGAGTGT TATTTTCATCCAGTTTGTCTC	IGF1	GATTTCTTGAAGGTGAAGATG GAAGGAAGTACATTTGAAGAA CAAGTAGAGGGAGTGCAGGAA CAAGAACTACAGGATGTAGGA CACAAATGCATGGGTGTTGTA	PPP2R2B	GACATTATCTCTACGGTAGAA CAGCTTACTTTCTTCTGTCTA GAGGATTAACCTATGGAACCT GAAATTATCTCTTCGATTTCCG TAATCTCACATACTGAATACT
AKT2	GACAAAGATGGCCACATCAAG GACTTTGGCCTCTGCAAAGAG GACACAAGGTACTTCGATGAT CAAGGTACTTCGATGATGAAT	IGF1R	CATCTTCGAGATGACCAATCT CATGTACTGCATCCCTTGTGA CATCTTACTACATGGGCTGAA GAGAATCCCAATGGATTGATT AAACACATTTGGGATGTTTCT	PTEN	GATGGATTTCGACTTAGACTTG AAGATCTTGACCAATGGCTAA TATTATAGTACCTGTTAAAG GATATCAAAGTAGAGTTCCTC CAATAGGACATTGTGTGATGAT
AKT3	AAAAACTGGAGGCCAAGATAC GAATGAATTGTAGTCCAACCT CACGTTTCTATGGTGCAGAAA GAAATGATGTGGGAGGTTA CATTCTGCTACTTCACTGTCA	INPP5D	GAATTGCGTTTACACTTACAG CATCAACATGGTGTCCAAGCT CATCTGTACTGACAACGTGAA GATTGAGTTTCTCAGTGCTA GATTTGAGGGTGGAGATAG	RPS6	GATGAACGCAAACTTCGTA GAAAATCAGTTCGTGGTTGCA AATCTGAGCGTTCTCAACTTG TACTTCTATGAGAAGCGTAT AAACTTCGTACTTTCTATGAG
AFX	GAAATCAGTCATATGCAGAAT CAGTTCAAATGCCAGCAGTGT CAAGACAGAATGCCTCAGGAT CATCAGTGACCTCATGGATGA CACTTAGGCTTTGTAGCAAGA	INSR	CATGGATATCCGGAACAACCT CACCAATACGTCATTACAAC CAGTATGCCATCTTTGTGAAG CACCTATTTCTACGTGACAGA GAGTTCAGAGATCGTTCCTAT	RPS6KA1	CATTTTCGAGCTCCTCAAGGTT TATCATTTACAGAGACCTCAA CATGTCTTCTACTCCACCATT GAAGAGATTGAGATTCTCTG AAAAATGGCATCAACCACCAT
CBL	GACAAGAAGATGGTGGAGAAG AAAAGACAATAGTCCCTTGGA GATAAGGATGTAAGATTGAG TATTCTGTTGGAGCAGAATCC GATATCACATCAGTGGTTCCA	IRS1	CACAAACGCTTCTTCGTA GATAATCGGTTCCGAAAGAGA CAGATCATCAATCCCATCAGA TACTCATTGCCAAGATCCTTT CAGAATGAAGACCTAAATGAC	RPS6KA3	GAGAATGGACAGCAAATTATG CAGAAGAAGATGTCAAATCT CAACGATAGACTGGAATAAAC AAAAGCAAGAGAGACCCAACA GATATTTGGATCCATGGTGA
CBLB	GATGCTATTCAGGATGCAGTT CAAAACTATCGTACCATGGAA TAATCCTGATTTAACTGGATT CAGAACTCACAGTACATCA TACATCTTAATGGTCAGAATT	mTOR	CACTATGTACCATGGAACCTC CAGATTCCACAGCTAAAGAAG GATGAAATAGTACCCTCATG GAGGTTATCCAGTACAACTT TAGTAAATGCTTCCACTAAAC	SGK	CAATTCATCGCTTTCATGA GAAAGGTTCTTCTAGCAAGAC CACTGAACATCGTTTTATAGAG GAAATGTACGACAACATTCTG TATGTGTGTTCCGAATGTTT
EIF4EBP1	GATCATCTATGACCGGAAATT CATCTATGACCGGAAATTCCT GAAGAGTACAGTTTGAGATG GACAAGAACGAACCCCTTCTT	PDK1	CAAAATCGGTTTCATTGATGA AAATGAAGATATGCACATCCA CAGAAGATCATTAAAGTTGGAA CAACATAGAGCAGTACATTCA GATAAGCGGAAGGTTTATTT	TSC1	TATGCTTGTAAACACCTTGGT TAGAAACTCATGATGTTGTGA AAAGAAGAAGCTGCAATATCT CATAATGCTGCCATGAAAGAT AAAAGGATAACCCAGGTGTTT
FKHR	GATCTACGAGTGGATGGTCAA AACAAACAGTAAATTTGCTAAG CAAAGATGGCCTCTACTTTAC TATACAAACACTTCAGGACAA TAAAAGTACTTCAGATTGTCT	PIK3CA	GAATGTTTACTACCAATGGA AAAAGTAGAAGTATGTTGCTA AAAATGGCTTTGAACTTTGG TACTGAAGAAAGCATTGACTA AATGAAAGCTCACTCTGGATT	TSC2	TAAACAGACGGAGTTTATCAT CATCAACAGGCAGTTCTATGA CAGCATTAACTCTTACCATA CAATGAGTCACAGTCCCTTGA CAGACATAGAGGCACAGATTG
FKHRL1	GACAATAGCAACAAGTATACC GATGAAGTCCAGGACGATGAT CATGTCACACTATGGTAACCA CATGTTCAATGGGAGCTTGGGA CAAACCTGACACAAGACCTACA	PIK3CB	GATTCACAGATAGCATCTGAT AAAACGTTTACTATTTCATG GACTTTAAAGGACAATTGAGA GATTGTGCCCTCTAGATTTC CATTGAGTGAACAGTAGCAA		
GSK3A	GACATCAAAGTATTGGCAAT CAAGTTGACCATCCCTATCCT CACTGATTACACCTCATCCAT CATTCTCATCCCTCCTCACTT GACTAGAGGGCAGAGTAAAT	PIK3R1	AAGAGAAGAAGATATTGACTT CAGAATTACGACAGCTTCTTG TACTGTAGCCAACAACGGTAT TACATGAATATAACAACCTCAGT GATCCTTCTCCTGAAGTTCAG		
GSK3B	CATGAAAGTTAGCAGAGACAA GAGAAAGCTAGATCACTGTAA CAATGTTTCGTATATCTGTTC GAGCAAATCAGAGAAATGAAC AAAACCACCAGTTACTTGAG	PIK3R2	CATTTCAAGGGAGGAGGTGAA CAATAAGCTGATCAAGGTCTT CAAATACCAGCAGGACCAGAT CAGATGAAGCGTACTGCAATT GAAAGAGATGCAAAGGATCCT		