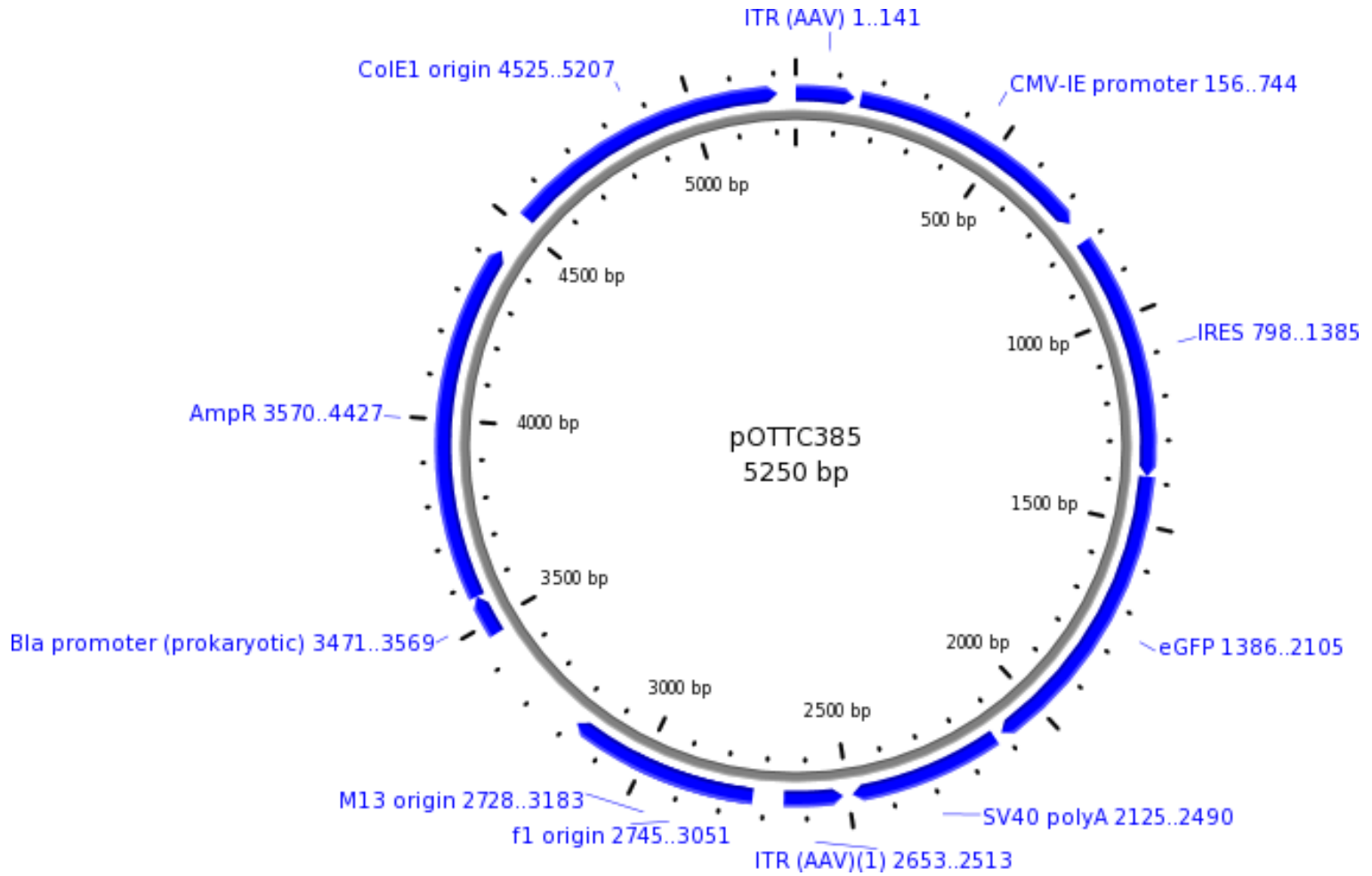


pAAV CMV IRES eGFP.dna.ape pOTTTC385



Name	Description	Location
ITR (AAV)	Inverted Terminal Repeat (Adeno-associated virus)	1..141
CMV-IE promoter	Human cytomegalovirus immediate early promoter	156..744
IRES	Internal Ribosomal Entry Site	798..1385
eGFP	Enhanced Green Fluorescent Protein	1386..2105
SV40 polyA	Simian virus 40 poly-adenylation signal	2125..2490
ITR (AAV)(1)	Inverted Terminal Repeat (Adeno-associated virus)	2653..2513
M13 origin	Phagemid origin of replication	2728..3183
f1 origin	Phagemid origin of replication	2745..3051
Bla promoter (prokaryotic)	Bacterial beta-lactamase promoter	3471..3569
AmpR	Ampicillin Resistance Marker	3570..4427
CoIE1 origin	Plasmid origin of replication	4525..5207

pOTTTC385 was constructed by Chris Richie at OTTC/NIDA (CR651_A).
 File dynamically generated on 23-JUL-2019 by NIDA plasmid manager.