

## 2021 2학기 과제 정리

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1) Goal : MGE(Mobile genetic element) & ARG(Antibiotic resistance gene) 둘 다 보유한 ORF(Open reading frame) 찾기

### 2) Resources

(1) CF1A-114.gff -> Contig & ORF 파일

```
##gff-version 3
##sequence-region k141_2 1 522
##sequence-region k141_4 1 621
##sequence-region k141_5 1 562
##sequence-region k141_7 1 511
##sequence-region k141_8 1 595
##sequence-region k141_9 1 686
##sequence-region k141_12 1 587
##sequence-region k141_13 1 755
##sequence-region k141_14 1 716
##sequence-region k141_15 1 840
##sequence-region k141_16 1 742
##sequence-region k141_17 1 967
##sequence-region k141_18 1 893
##sequence-region k141_23 1 555
##sequence-region k141_25 1 525
##sequence-region k141_29 1 673
##sequence-region k141_30 1 1247
##sequence-region k141_31 1 574
##sequence-region k141_32 1 607
##sequence-region k141_33 1 618
##sequence-region k141_35 1 704
##sequence-region k141_36 1 837
##sequence-region k141_37 1 522
##sequence-region k141_38 1 1608
##sequence-region k141_39 1 508
##sequence-region k141_40 1 545
##sequence-region k141_41 1 599
##sequence-region k141_42 1 554
##sequence-region k141_43 1 605
##sequence-region k141_44 1 584
##sequence-region k141_45 1 502
##sequence-region k141_47 1 1685
##sequence-region k141_48 1 713
##sequence-region k141_49 1 1957
##sequence-region k141_50 1 1014
##sequence-region k141_51 1 594
##sequence-region k141_52 1 591
##sequence-region k141_53 1 686
```

##sequence  
필요 없는 것

/CDS 입력시

필요한 영역으로 바로 이동  
(Contig-ORF)

Contig name

ORF name

Contig name	ORF name
k141_2	Prodigal:2.6 CDS 49 477 . - 0 ID=ALLJNDBJ_00001;inference=ab initio prediction:Prodigal:2.6;locus_tag=ALLJNDBJ_00001;product=hypothetical protein
k141_4	Prodigal:2.6 CDS 7 255 . - 0 ID=ALLJNDBJ_00002;inference=ab initio prediction:Prodigal:2.6;locus_tag=ALLJNDBJ_00002;product=hypothetical protein
k141_4	Prodigal:2.6 CDS 256 459 . - 0 ID=ALLJNDBJ_00003;inference=ab initio prediction:Prodigal:2.6;locus_tag=ALLJNDBJ_00003;product=hypothetical protein
k141_5	Prodigal:2.6 CDS 136 465 . + 0 ID=ALLJNDBJ_00004;inference=ab initio prediction:Prodigal:2.6;locus_tag=ALLJNDBJ_00004;product=hypothetical protein
k141_9	Prodigal:2.6 CDS 42 296 . + 0 ID=ALLJNDBJ_00005;inference=ab initio prediction:Prodigal:2.6;locus_tag=ALLJNDBJ_00005;product=hypothetical protein
k141_12	Prodigal:2.6 CDS 250 555 . - 0 ID=ALLJNDBJ_00006;inference=ab initio prediction:Prodigal:2.6;locus_tag=ALLJNDBJ_00006;product=hypothetical protein
k141_15	Prodigal:2.6 CDS 46 216 . + 0 ID=ALLJNDBJ_00007;inference=ab initio prediction:Prodigal:2.6;locus_tag=ALLJNDBJ_00007;product=hypothetical protein
k141_16	Prodigal:2.6 CDS 148 723 . - 0 ID=ALLJNDBJ_00008;inference=ab initio prediction:Prodigal:2.6;locus_tag=ALLJNDBJ_00008;product=hypothetical protein
k141_17	Prodigal:2.6 CDS 347 469 . - 0 ID=ALLJNDBJ_00009;inference=ab initio prediction:Prodigal:2.6;locus_tag=ALLJNDBJ_00009;product=hypothetical protein
k141_18	Prodigal:2.6 CDS 7 834 . + 0 ID=ALLJNDBJ_00010;inference=ab initio prediction:Prodigal:2.6;locus_tag=ALLJNDBJ_00010;product=hypothetical protein
k141_30	Prodigal:2.6 CDS 56 964 . + 0 ID=ALLJNDBJ_00011;inference=ab initio prediction:Prodigal:2.6;locus_tag=ALLJNDBJ_00011;product=hypothetical protein
k141_36	Prodigal:2.6 CDS 282 833 . - 0 ID=ALLJNDBJ_00012;inference=ab initio prediction:Prodigal:2.6;locus_tag=ALLJNDBJ_00012;product=hypothetical protein

```
k141_664575 Prodigal:2.6 CDS 367 768 . -
similar to AA sequence:UniProtKB:P32055;locus tag=ALLJNDBJ 730201;
##FASTA
>k141_2
GCCGGGCGGATTCGTTCCGCCAGGCGGCGGAGCAGGCCAGCTGACCTTAAACTCCGCG
GCATCCGTGATCCGGGAGCAGTTTGCAGGGGATTCCATCGAACTGGTAAACACCTACACC
ACCGTGACCAACACCTCCGGAGGCATCACCGGTGACCAAGAATCCCGGAACAGTGACG
GTTTCTACAGCAATTCCAACGGCAAGGGACAGGAAACGGCTCTCGCCTCCGGCACCTAT
TCCCAGGCAAGCGGATTGAATCTGATCCAGGGAAGTCTGCCACAGCTGCGCAGGGGACTG
CTGAATTGGGCGATTGACACCATGACAGGCATCGTGCTGACCAATAATGAATGCTCGGCC
AACTATGTGGTCAAAGCCAATGGGCCGGAGGGCGCTCTGGAGGATGTGCGGGTGAAGATT
CGCATGGAGCCCGCGGCCACAGCGGATTGGCCACCCAGGATGAAAGGCAGTCCCATGAT
GCGGAGAAGTACTACATGACTGTTTTGTTCTCCCTGGAGTCC
>k141_4
CTATCCCTATTGGTGAAGTCCACAAGTTCGCCGTACGCAAGAACGAAACCCGCATCGCA
CGCATGCAGCGCCTGGGCAACGACAAGCGGACGGCATCTTCCCGACAAGATGGTGGA
GAAAACAAGTTCTGCTGCACTGGCCGACAACAAGGAGCTGGGAGCCTACATGGAGCAG
AAAAAGGAATGGATCGAGGAGGAACCCCTTCGTCAAGAACTCTACAACACACTCATCGAA
AGCGACATCTTCCAATATACCTGACCAAGGAGGAATTCGACTATGAAGCCGACCGCGAG
CTGGTACGGAAGTTCTACAAGACGTACGTCTGCAACAACGAAGGTGTGGAAGACCTGATC
GAGGACCACTGCCTCTACTGGAACGACGACCGCTTCGTGCTGCACTCCTTCGTCTGAAG
ACGATAAAGCGCTTCGCGCAGGCCGCGCAGCGACCAACCGCTGCTGCGCGCAGTTTGCC
AACGAAGAGGACCGCGAGTTTGCCGCAAACTCTTCGCCGACGCCATCAATAACGAGTCC
CGCACCCGCATCATCATCCGCGAAACTGCAAGAAGTGGGAGTTTGACCGACTTGCTTTC
ATGGACGTCATCATCATGACG
>k141_5
:
```

##FASTA

이후에는 필요 없는 것

즉 전체에서  
##sequence  
필요한 것  
##FASTA

- \* 하나의 contig안에 여러개의 ORF 있을 수 있음
- \* /CDS, /##FASTA처럼 입력 해 원하는 위치로 이동

\*\* 전체에서 ##sequence ~ ##FASTA 사이 contig - orf name 나온 것만 필요함



(2) CF1A-114.f.dia -> ORF & ARG 파일

ORF name		ARG	
ALLJNDBJ_10202	gb A_G06942.1 ARO:300298	arnA	71.4 70 20 0 2 71 582 651 1.4e-25 108.6
ALLJNDBJ_15370	gb A_C75089.1 ARO:300357	ugd	70.2 198 59 0 4 201 191 388 5.2e-82 297.4
ALLJNDBJ_16978	gb B_B38260.1 ARO:300083	cpxA	100.0 171 0 0 1 171 287 457 5.8e-98 350.1
ALLJNDBJ_24130	gb A_A88675.1 ARO:300049	ErmF	95.3 150 7 0 1 150 117 266 4.8e-80 290.4
ALLJNDBJ_27880	gb B_D59497.1 ARO:300050	Nocardia	77.7 130 28 1 1 130 1009 1137 2.2e-55 208.8
ALLJNDBJ_38892	gb B_B36671.1 ARO:300083	evgA	100.0 204 0 0 1 204 1 204 7.6e-113 399.8
ALLJNDBJ_41860	gb C_H51823.1 ARO:300055	tet44	98.9 640 7 0 1 640 1 640 0.0e+00 1265.4
ALLJNDBJ_41861	gb C_H51824.1 ARO:300262	ANT(6)-Ib	100.0 285 0 0 1 285 1 285 2.5e-170 591.3
ALLJNDBJ_53447	gb A_A71728.1 ARO:300293	vanSG	77.0 61 14 0 2 62 305 365 1.3e-21 95.1
ALLJNDBJ_56153	gb C_A79727.1 ARO:300019	tetQ	90.0 330 33 0 3 332 328 657 3.3e-174 604.4
ALLJNDBJ_59366	gb A_V10830.1 ARO:300264	APH(3')-IIa	100.0 264 0 0 1 264 1 264 1.7e-157 548.5
ALLJNDBJ_60536	gb A_A23018.1 ARO:300445	Campylobacter	100.0 207 0 0 1 207 1 207 9.4e-127 446.0
ALLJNDBJ_65471	gb C_G86592.1 ARO:300462	Erm(49)	100.0 276 0 0 1 276 1 276 9.4e-160 556.2
ALLJNDBJ_70282	gb A_A26652.1 ARO:300283	lnuA	95.9 161 5 0 1 161 1 161 1.8e-93 335.1
ALLJNDBJ_82405	gb A_U10334.1 ARO:300262	aad(6)	99.3 135 1 0 1 135 142 276 1.5e-77 282.0
ALLJNDBJ_82406	gb A_B53445.1 ARO:300289	SAT-4	99.4 180 1 0 1 180 1 180 5.7e-96 343.6
ALLJNDBJ_88596	gb C_A79727.1 ARO:300019	tetQ	94.9 79 4 0 1 79 579 657 6.7e-41 159.5
ALLJNDBJ_90209	gb C_A79727.1 ARO:300019	tetQ	98.7 239 3 0 1 239 419 657 3.6e-138 484.2
ALLJNDBJ_91147	gb A_C75089.1 ARO:300357	ugd	78.5 200 53 0 2 201 189 388 1.2e-86 312.8
ALLJNDBJ_95690	gb A_C75271.1 ARO:300395	yojI	100.0 114 0 0 1 114 434 547 2.4e-60 224.6
ALLJNDBJ_108035	gb A_D23513.1 ARO:300300	CfxA2	78.6 318 68 0 1 318 1 318 1.2e-141 496.1
ALLJNDBJ_113165	gb C_I79240.1 ARO:300504	leptB	85.9 61 8 0 1 61 510 570 4.1e-29 120.2
ALLJNDBJ_121197	gb B_A16547.1 ARO:300002	emrA	99.0 204 2 0 1 204 187 390 2.4e-111 394.8
ALLJNDBJ_125784	gb C_A26199.1 ARO:300408	ANT(3')-IIa	100.0 266 0 0 48 313 58 323 4.8e-151 527.3
ALLJNDBJ_130198	gb A_P42147.1 ARO:300444	tet(W/M/W)	92.2 639 50 0 1 639 1 639 0.0e+00 1175.2
ALLJNDBJ_137566	gb A_A20117.1 ARO:300019	tetB(P)	98.0 201 4 0 1 201 333 533 4.5e-110 390.6
ALLJNDBJ_137567	gb A_A20117.1 ARO:300019	tetB(P)	100.0 84 0 0 1 84 569 652 1.4e-44 171.8
ALLJNDBJ_141755	gb B_E78084.1 ARO:300354	mdtN	99.7 293 1 0 1 293 51 343 1.9e-157 548.5
ALLJNDBJ_145918	gb B_F80809.1 ARO:300266	npmA	100.0 219 0 0 1 219 1 219 2.3e-123 434.9
ALLJNDBJ_150933	gb A_A23033.2 ARO:300019	tetO	95.9 98 4 0 1 98 294 391 2.6e-48 184.9
ALLJNDBJ_155028	gb A_L79549.2 ARO:300300	CfxA3	92.8 319 23 0 52 370 3 321 3.6e-169 587.8
ALLJNDBJ_157745	gb A_H87088.1 ARO:300019	tet32	78.9 594 155 0 1 594 46 639 1.0e-263 902.5
ALLJNDBJ_166187	gb A_C23556.1 ARO:300266	APH(6)-Id	99.0 209 2 0 1 209 70 278 3.6e-118 417.5
ALLJNDBJ_170473	gb A_N06707.1 ARO:300016	tet(A)	99.7 390 1 0 1 390 26 415 1.2e-218 752.3
ALLJNDBJ_171969	gb B_A12910.1 ARO:300031	mphB	85.4 295 43 0 1 295 5 299 2.4e-152 531.6
ALLJNDBJ_178705	gb A_P42147.1 ARO:300444	tet(W/M/W)	93.8 160 10 0 2 161 479 638 7.9e-89 319.7
ALLJNDBJ_196886	gb C_M12479.1 ARO:300056	tet(40)	97.9 382 8 0 1 382 25 406 1.2e-199 689.1
ALLJNDBJ_197078	gb B_A12910.1 ARO:300031	mphB	84.9 86 13 0 1 86 214 299 2.2e-40 157.9
ALLJNDBJ_214036	gb C_A79727.1 ARO:300019	tetQ	98.8 113 7 0 1 113 545 657 3.9e-58 217.2
ALLJNDBJ_218583	gb A_Y32951.1 ARO:300283	lnuC	88.4 164 19 0 1 164 1 164 1.2e-87 315.8
ALLJNDBJ_218629	gb A_F74725.1 ARO:300465	Mef(En2)	99.3 401 3 0 1 401 1 401 5.7e-221 760.0
ALLJNDBJ_221421	gb C_E51638.1 ARO:300264	APH(3')-Ia	98.2 271 5 0 1 271 1 271 4.8e-163 567.0
ALLJNDBJ_227090	gb A_I44920.1 ARO:300447	poxTA	76.7 536 125 0 1 536 1 536 1.9e-240 825.1

(3) CF1A-114.f.Int.dia -> ORF & MGE(Int)

ORF name		MGE											
ALLJNDBJ_36051	gi 03466236 ref WP_013700897.1	97.3	182	5	0	1	182	101	282	1.6e-97	349.0	182	410
ALLJNDBJ_83395	gi 79963920 ref WP_045411042.1	70.3	138	41	0	2	139	277	414	6.4e-57	213.8	143	421
ALLJNDBJ_107001	gi 92635861 gb EFF54355.1	76.8	267	62	0	1	267	19	285	1.4e-121	429.5	267	285
ALLJNDBJ_146474	GCA_000012825.1_CP000139.1.chr.fa_1_1	85.5	62	9	0	1	62	206	267	1.7e-27	114.8	62	267
ALLJNDBJ_266196	GCA_000325705.1_CP003346.1.chr.fa_2_3	71.6	109	31	0	1	109	199	307	3.7e-44	171.0	111	308
ALLJNDBJ_327045	GCA_000325705.1_CP003346.1.chr.fa_2_3	71.8	103	29	0	1	103	206	308	2.4e-42	164.9	103	308
ALLJNDBJ_328113	gi 95296399 ref WP_008021152.1	73.5	219	58	0	1	219	49	267	2.8e-96	345.1	219	267
ALLJNDBJ_350508	Int_1_gi_394348876_gb_CP003684.1_392	100.0	337	0	0	1	337	1	337	2.1e-199	688.3	337	337
ALLJNDBJ_360151	GCA_000012825.1_CP000139.1.chr.fa_1_1	100.0	267	0	0	1	267	1	267	9.5e-155	539.7	267	267
ALLJNDBJ_504227	Int_1_gi_516560780_gb_JX515588.1_219	70.9	55	16	0	16	70	237	291	2.2e-18	84.7	70	291
ALLJNDBJ_507147	gi 03466236 ref WP_013700897.1	100.0	74	0	0	1	74	337	410	6.9e-39	152.9	74	410
ALLJNDBJ_507148	gi 03466236 ref WP_013700897.1	97.8	91	2	0	1	91	231	321	1.7e-50	191.8	92	410
ALLJNDBJ_545043	gi 85780171 ref WP_001403201.1	97.0	233	7	0	1	233	189	421	1.8e-133	468.8	233	421
ALLJNDBJ_616397	gi 24765482 emb CDE64103.1	74.4	156	40	0	2	157	112	267	1.2e-69	256.1	157	267
ALLJNDBJ_660926	gi 95115106 ref WP_007839924.1	71.7	219	62	0	1	219	49	267	1.2e-94	339.7	219	267
ALLJNDBJ_668589	gi 46399651 ref WP_039441885.1	72.0	264	74	0	6	269	4	267	2.7e-117	415.2	269	267
(END)													

(4) CF1A-114.f.IS.dia -> ORF & MGE(IS)

ORF name		MGE												
ALLJNDBJ_02745	ISLj05	70.2	131	39	0	1	131	300	430	1.8e-52	199.9	139	445	
ALLJNDBJ_06165	ISCce2	85.4	89	13	0	7	95	305	393	1.3e-41	163.3	97	398	
ALLJNDBJ_06211	ISCth10	75.9	137	33	0	1	137	19	155	3.4e-59	222.2	138	158	
ALLJNDBJ_06763	ISCth10	78.7	122	26	0	1	122	1	122	7.9e-56	211.1	140	158	
ALLJNDBJ_10662	IS3411	89.7	107	11	0	1	107	1	107	3.3e-51	195.7	136	390	
ALLJNDBJ_10663	ISEc39	87.7	81	7	1	1	78	1	81	2.3e-33	135.6	78	402	
ALLJNDBJ_12359	ISEc12	99.1	111	1	0	1	111	139	249	4.2e-60	224.9	111	249	
ALLJNDBJ_14136	ISFnu8	71.4	112	32	0	1	112	339	450	2.6e-41	162.5	114	454	
ALLJNDBJ_15856	ISLjo1	84.0	125	20	0	15	139	225	349	2.3e-60	226.1	139	349	
ALLJNDBJ_16715	ISBaov1	100.0	409	0	0	20	428	1	409	1.0e-247	850.1	428	409	
ALLJNDBJ_18390	ISBf13	80.2	86	17	0	1	86	38	123	3.6e-35	141.7	86	367	
ALLJNDBJ_20236	ISRgn1	79.0	200	42	0	1	200	185	384	1.0e-93	337.4	201	386	
ALLJNDBJ_20832	ISG12	99.2	240	2	0	1	240	190	429	5.5e-142	498.0	240	429	
ALLJNDBJ_24486	ISBf11	92.3	117	9	0	2	118	280	396	5.9e-60	224.6	121	428	
ALLJNDBJ_32627	ISBco10	70.0	150	44	1	2	150	204	353	1.1e-55	210.7	155	470	
ALLJNDBJ_34267	ISEnfa1	0	74.1	220	56	1	1	220	182	400	6.6e-89	321.6	222	402
ALLJNDBJ_34930	ISBco9	75.0	96	24	0	1	96	280	375	2.9e-41	162.2	98	376	
ALLJNDBJ_38985	ISLhe6	94.7	38	2	0	1	38	1	38	9.1e-15	72.8	38	411	
ALLJNDBJ_38986	ISLhe6	86.8	53	7	0	1	53	46	98	1.5e-20	92.4	53	411	
ALLJNDBJ_40276	ISStrsp	73.1	245	65	1	29	272	216	460	1.8e-107		383.6	280	461
ALLJNDBJ_46758	ISG14	97.7	218	5	0	1	218	212	429	4.6e-127	448.4	218	429	
ALLJNDBJ_47950	ISCpe2	76.4	89	21	0	1	89	296	384	8.0e-38	150.6	89	384	
ALLJNDBJ_50041	ISCce2	89.9	149	15	0	1	149	105	253	1.9e-79	289.7	149	398	
ALLJNDBJ_51156	IS200F	73.3	101	27	0	1	101	41	141	3.0e-44	172.2	104	152	
ALLJNDBJ_51677	ISG29	95.6	295	13	0	1	295	110	404	2.6e-165	575.9	295	404	
ALLJNDBJ_51678	ISG29	97.2	108	3	0	1	108	1	108	1.0e-55	210.3	108	404	
ALLJNDBJ_57524	ISSpn6	72.0	50	14	0	1	50	89	138	1.8e-18	85.5	50	157	
ALLJNDBJ_58953	ISBas1	73.6	485	127	1	1	485	1	484	1.5e-207	716.8	485	484	
ALLJNDBJ_60630	ISG09	99.0	96	1	0	1	96	307	402	1.9e-45	176.0	96	402	
ALLJNDBJ_60810	ISStin1	77.9	149	33	0	1	149	1	149	7.1e-67	248.1	172	154	
ALLJNDBJ_61939	ISLhe65	76.5	388	89	1	2	387	1	388	2.2e-175	609.8	399	395	
ALLJNDBJ_62328	ISBian1	82.3	141	25	0	5	145	1	141	1.4e-63	236.9	145	321	
ALLJNDBJ_63220	IS1201	94.6	147	8	0	1	147	120	266	1.9e-79	289.7	152	369	
ALLJNDBJ_63260	ISLgar1	91.5	201	17	0	1	201	248	448	6.6e-104	371.3	202	449	
ALLJNDBJ_66374	ISG13	89.3	428	46	0	1	428	1	428	2.2e-234	805.8	428	428	
ALLJNDBJ_66832	ISStrsp	71.8	262	74	0	1	262	198	450	1.5e-111		397.1	264	461
ALLJNDBJ_67102	ISLhe4	98.8	409	5	0	1	409	1	409	3.2e-206	712.2	409	409	
ALLJNDBJ_67724	IS15410	72.2	151	41	1	4	154	3	152	1.0e-64	240.7	154	152	
ALLJNDBJ_68064	ISCce2	77.3	238	52	1	1	236	161	398	7.7e-104	371.3	236	398	
ALLJNDBJ_72212	ISDha13	73.8	145	38	0	6	150	7	151	6.1e-62	231.5	153	151	
ALLJNDBJ_72791	ISLjo5	77.0	81	18	0	1	81	362	442	1.3e-37	149.8	83	445	
ALLJNDBJ_73246	ISBco2	81.6	250	46	0	1	250	1	250	8.9e-119	421.0	250	250	
ALLJNDBJ_73304	IS100ky	100.0	220	0	0	0	1	220	40	259	1.7e-121	429.9	220	259

3) test01.py -> 추출

(1) import, 필요한 자료 구조 생성

```
import sys
import pandas as pd
from Bio.SeqIO.FastaIO import SimpleFastaParser as SFP

M_Int = sys.argv[1]
M_IS = sys.argv[2]
Arg = sys.argv[3]
Ctg = sys.argv[4]
new = sys.argv[5]

M_Int_ORF_list = []
M_Int_Name_list = []
M_IS_ORF_list = []
M_IS_Name_list = []
Arg_ORF_list = []
Arg_Name_list = []
Ctg_CTG_list = []
Ctg_ORF_list = []
```

(2) MGE, ARG에서 필요한 부분만 받아오기

```
with open(M_Int, 'r') as M_Int, open(M_IS, 'r') as M_IS, open(Arg, 'r') as Arg:
    for line in M_Int:
        line = line.split()
        M_Int_orf = line[0]
        M_Int_name = line[1]
        M_Int_ORF_list.append(M_Int_orf)
        M_Int_Name_list.append(M_Int_name)
    for line in M_IS:
        line = line.split()
        M_IS_orf = line[0]
        M_IS_name = line[1].split('|')[0]
        M_IS_ORF_list.append(M_IS_orf)
        M_IS_Name_list.append(M_IS_name)
    for line in Arg:
        line = line.split()
        Arg_orf = line[0]
        Arg_name = line[1].split('|')[3]
        Arg_ORF_list.append(Arg_orf)
        Arg_Name_list.append(Arg_name)

M_Int_df = pd.DataFrame({'ORF':M_Int_ORF_list, 'Name':M_Int_Name_list})
M_IS_df = pd.DataFrame({'ORF':M_IS_ORF_list, 'Name':M_IS_Name_list})
Arg_df = pd.DataFrame({'ORF':Arg_ORF_list, 'Name':Arg_Name_list})
M_df = M_Int_df.append(M_IS_df, ignore_index = True)
#print(M_Int_df)
#print(M_IS_df)
#print(M_df)
#print(Arg_df)
```

필요한 ORF,  
GENE NAME 만  
받기

Dataframe에  
옮기기

ex ) M\_IS (\*\*M\_Int 파일이어야 하는데... 잘못 적은 듯..)

① line = line.split() -> 라인마다 공백으로 split

ALLJNDBJ\_36051 | gi|503466236|ref|WP\_013700897.1| 97.3 182 5 0 1 182 101 282 1.6e-97 349.0 182 410

② M\_IS\_orf = line[0]

ALLJNDBJ\_36051

③ M\_IS\_name = line[1].split('|')[0]

gi|503466236|ref|WP\_013700897.1|

gi

④ M\_IS\_ORF\_list.append(M\_IS\_orf) => ORF 어펜드

⑤ M\_IS\_NAME\_list.append(M\_IS\_name) => MGE name 어펜드



### (3) Contig-ORF 추출

```
with open(Ctg, 'r') as Ctg:
    for line in Ctg:
        if line.startswith('##g') | line.startswith('##s'):
            pass
        elif line.startswith('##F'):
            break
        else:
            line = line.split()
            Ctg_ctg = line[0]
            Ctg_orf = line[8].split('=')[1].split(';')[0]
            Ctg_CTG_list.append(Ctg_ctg)
            Ctg_ORF_list.append(Ctg_orf)

Ctg_df = pd.DataFrame({'CTG':Ctg_CTG_list, 'ORF':Ctg_ORF_list})
#print(Ctg_df)
```

① '##g', '##s'로 시작하면 pass

```
##gff-version 3
##sequence-region k141_2 1 522
##sequence-region k141_4 1 621
##sequence-region k141_5 1 562
##sequence-region k141_7 1 511
##sequence-region k141_8 1 595
##sequence-region k141_9 1 686
```

첫 부분(필요 x)  
pass

② '##F'로 시작하면 break

```
##FASTA
>k141_2
GCCGGGCGGATTCGTTCCGCCAGGCGGCGGAGCAGGCCAGCTGACCTTAAACTCCGCG
GCATCCGTGATCCGGGAGCAGTTTGCAGGGGATTCCATCGAACTGGTAAACACCTACACC
ACCGTGACCAACACCTCCGGAGGCATCACCACGGTGACCAAGAATCCCGGAACAGTGACG
GTTTCCTACAGCAATTCCAACGGCAAGGGACAGGAAACGGCTCTCGCCTCCGGCACCTAT
TCCAGGCAAGCGGATTGAATCTGATCCAGGGAAGTCTGCCACAGCTGCGCAGGGGACTG
CTGAATTGGGCGATTGACACCATGACAGGCATCGTGCTGACCAATAATGAATGCTCGGCC
AACTATGTGGTCAAAGCCAATGGGCCGGAGGGCGCTCTGGAGGATGTGCGGGTGAAGATT
CGCATGGAGCCCGGCCACAGCGGATTTGGCCACCCAGGATGAAAGGCAGTCCCATGAT
GCGGAGAAGTACTACATGACTGTTTTGTTCTCCCTGGAAGTCC
>k141_4
```

끝 부분(필요 x)  
더 이상 읽을 필요 x  
break

③ line = line.split(), Ctg\_ctg = line[0], Ctg\_orf = line[8].split('=')[1].split(';')[0]

k141_2	Prodigal:2.6	CDS	49	477	.	.	0	ID=ALLJNDBJ_00001;inference=ab initio prediction:Prodigal:2.6;locus_tag=ALLJNDBJ_00001;product=hypothetical protein
k141_4	Prodigal:2.6	CDS	7	255	.	.	0	ID=ALLJNDBJ_00002;inference=ab initio prediction:Prodigal:2.6;locus_tag=ALLJNDBJ_00002;product=hypothetical protein
k141_4	Prodigal:2.6	CDS	256	459	.	.	0	ID=ALLJNDBJ_00003;inference=ab initio prediction:Prodigal:2.6;locus_tag=ALLJNDBJ_00003;product=hypothetical protein

Ctg\_ctg = line[0]

Ctg\_orf = line[8]

```
line[8].split('=')[1]
```

```
ID=ALLJNDBJ_00001;inference=ab initio prediction:Prodigal:2.6;locus_tag=ALLJNDBJ_00001;product=hypothetical protein
ID=ALLJNDBJ_00002;inference=ab initio prediction:Prodigal:2.6;locus_tag=ALLJNDBJ_00002;product=hypothetical protein
ID=ALLJNDBJ_00003;inference=ab initio prediction:Prodigal:2.6;locus_tag=ALLJNDBJ_00003;product=hypothetical protein
```

```
line[8].split('=')[1].split(':')[0]
```

```
=ALLJNDBJ_00001;inference=
=ALLJNDBJ_00002;inference=
=ALLJNDBJ_00003;inference=
```

```
=ALLJNDBJ_00001
=ALLJNDBJ_00002
=ALLJNDBJ_00003
```

결과물  
->ORF만 추출

(4) merge

① 현재까지 생성된 데이터 프레임 3개

M\_df

ORF	Name
.	.
.	.
.	.

Arg\_df

ORF	Name
.	.
.	.
.	.

Ctg\_df

CTG	ORF
.	.
.	.
.	.

② 데이터 프레임 합치기

```
M_C_df = pd.merge(left = M_df, right = Ctg_df, how = 'left', left_on = 'ORF', right_on = 'ORF')
A_C_df = pd.merge(left = Arg_df, right = Ctg_df, how = 'left', left_on = 'ORF', right_on = 'ORF')
```

=> M\_df + Ctg\_df = M\_C\_df

ORF	Name
.	.
.	.
.	.

+

CTG	ORF
.	.
.	.
.	.

=

M	ORF	CTG
.	.	.
.	.	.
.	.	.

how = 'left' -> 왼쪽 데이터 프레임(M\_df)을 기준으로 병합  
\*\* 두 데이터 프레임은 ORF열을 통해 병합됨  
\*\* 왼쪽이 기준으로 MGE가 있는 ORF에 해당하는 contig만 알아냄

### ③ Kind 적기

```
M_C_df['Kind'] = 'MGE'  
A_C_df['Kind'] = 'ARG'
```

### (4) list comprehension

```
M_C_li = M_C_df['CTG'].tolist()  
A_C_li = A_C_df['CTG'].tolist()  
M_A_C_li = [x for x in A_C_li if x in M_C_li]
```

M\_A\_C\_li = [x for x in A\_C\_li if x in M\_C\_li]

=> 만약 A\_C\_li의 x가 M\_C\_li에도 있으면 x를 M\_A\_C\_li 리스트에 삽입한다.

\*\* A\_C\_li가 M\_C\_li보다 양이 많기 때문에 A\_C\_li의 x로 설정

### (5) 최종 + 저장

```
last_M_df = M_C_df.loc[M_C_df['CTG'].isin(M_A_C_li)]  
last_A_df = A_C_df.loc[A_C_df['CTG'].isin(M_A_C_li)]  
  
#print(last_M_df)  
#print(last_A_df)  
  
last_df = last_M_df.append(last_A_df)  
last_df = last_df.sort_values(by=['CTG', 'ORF'])  
last_df = last_df.reset_index()  
last_df = last_df[['CTG', 'ORF', 'Name', 'Kind']]  
  
#print(last_df)  
  
#pd.DataFrame(df).fillna('0').to_csv(sys.argv[1]+'total_re.csv',sep=',')  
last_df.to_csv(new)
```

\*\* 참고 : 진주언니가 만든 파일 -> 3inc\_ctg.py



#### 4) find\_ctg.sh

```
#M_Int = sys.argv[1] #/home/bbang9/Project/2020/CDC/20_03/HiSeq/ARG_finding/Integrase/filtered
#M_IS = sys.argv[2] #/home/bbang9/Project/2020/CDC/20_03/HiSeq/ARG_finding/IS/filtered
#Arg = sys.argv[3] #/home/bbang9/Project/2020/CDC/20_03/HiSeq/ARG_finding/Contig/filtered
#Ctg = sys.argv[4] #/home/bbang9/Project/2020/CDC/20_03/HiSeq/EDGE/CF1A-114/prokka_annot
#new = sys.argv[5]

gffpath='/home/bbang9/Project/2020/CDC/20_03/HiSeq/EDGE/'
argmge='/home/bbang9/Project/2020/CDC/20_03/HiSeq/ARG_finding/'
try='/home/guest01/2021/yb/yb01/Jinju/test01.py'
output='/home/guest01/2021/yb/yb01/Jinju/out/'
```

#### 패스 설정

```
#python ${try} M_Int M_IS ARG Ctg

for sample in $(gffpath)*
do
# echo ${sample}
ID=$(sample#gffpath)
python ${try} $(argmge)Integrase/filtered/${ID}.f.dia $(argmge)IS/filtered/${ID}.f.dia $(argmge)Contig/filtered/${ID}.f.dia $(gffpath)${ID}/prokka_annot/${ID}.gff ${output}${ID}.mobilome.out
done
```

#### 리눅스 쉘 스크립트

#### find\_ctg.sh

#### 5) output

```
[guest01@smel0:Jinju]$ cd out
[guest01@smel0:out]$ ll
total 296
-rw-rw-r-- 1 guest01 guest01 894 Sep 30 13:09 CF1A-114.mobilome.out
-rw-rw-r-- 1 guest01 guest01 743 Sep 30 13:09 CF1A-124.mobilome.out
-rw-rw-r-- 1 guest01 guest01 991 Sep 30 13:09 CF1A-1314.mobilome.out
-rw-rw-r-- 1 guest01 guest01 745 Sep 30 13:09 CF1A-1324.mobilome.out
-rw-rw-r-- 1 guest01 guest01 1233 Sep 30 13:09 CF1A-1334.mobilome.out
-rw-rw-r-- 1 guest01 guest01 803 Sep 30 13:09 CF1A-134.mobilome.out
-rw-rw-r-- 1 guest01 guest01 945 Sep 30 13:09 CF1A-314.mobilome.out
-rw-rw-r-- 1 guest01 guest01 1376 Sep 30 13:09 CF1A-324.mobilome.out
-rw-rw-r-- 1 guest01 guest01 891 Sep 30 13:09 CF1A-3314.mobilome.out
-rw-rw-r-- 1 guest01 guest01 761 Sep 30 13:09 CF1A-3324.mobilome.out
-rw-rw-r-- 1 guest01 guest01 338 Sep 30 13:09 CF1A-3334.mobilome.out
-rw-rw-r-- 1 guest01 guest01 990 Sep 30 13:09 CF1A-334.mobilome.out
-rw-rw-r-- 1 guest01 guest01 854 Sep 30 13:09 CF1E-111.mobilome.out
-rw-rw-r-- 1 guest01 guest01 1207 Sep 30 13:09 CF1E-112B.mobilome.out
-rw-rw-r-- 1 guest01 guest01 1603 Sep 30 13:09 CF1E-112P.mobilome.out
-rw-rw-r-- 1 guest01 guest01 1360 Sep 30 13:09 CF1E-311.mobilome.out
-rw-rw-r-- 1 guest01 guest01 1349 Sep 30 13:09 CF1E-312B.mobilome.out
-rw-rw-r-- 1 guest01 guest01 1227 Sep 30 13:09 CF1E-312P.mobilome.out
-rw-rw-r-- 1 guest01 guest01 837 Sep 30 13:09 CF1H-114.mobilome.out
```