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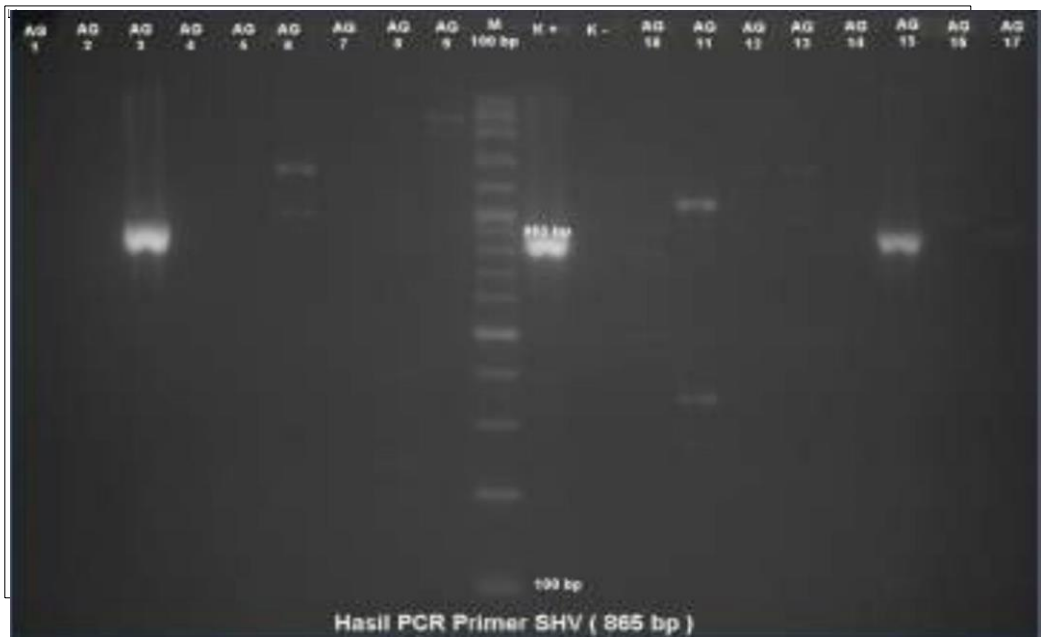
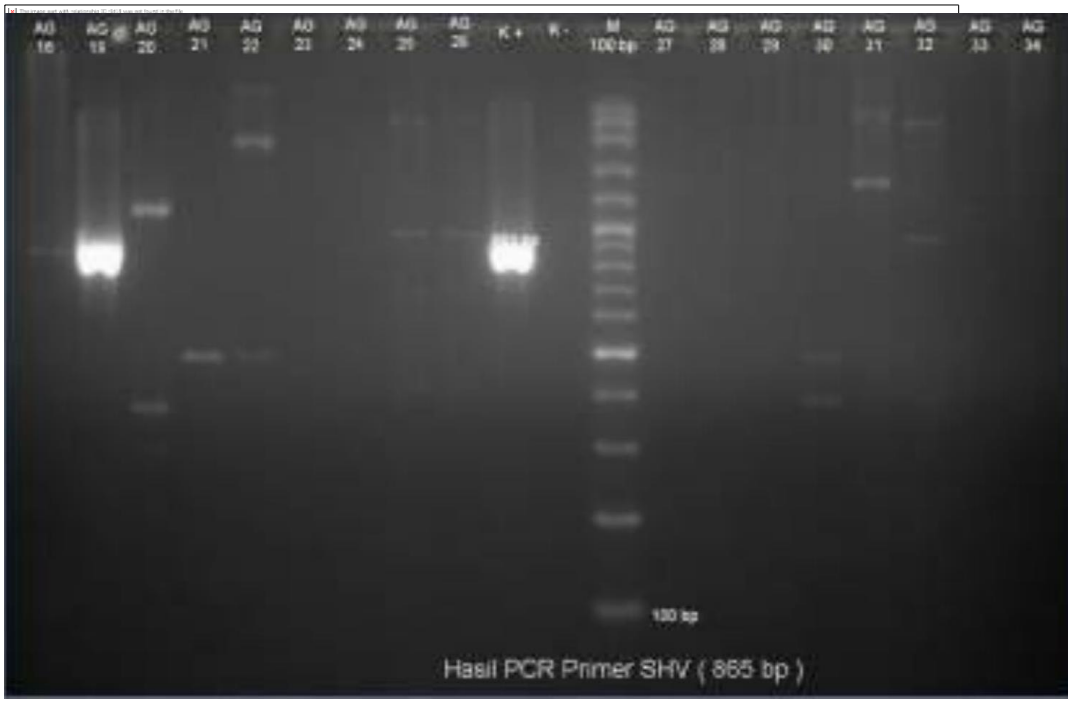
Lampiran 1. Hasil elektroforesis

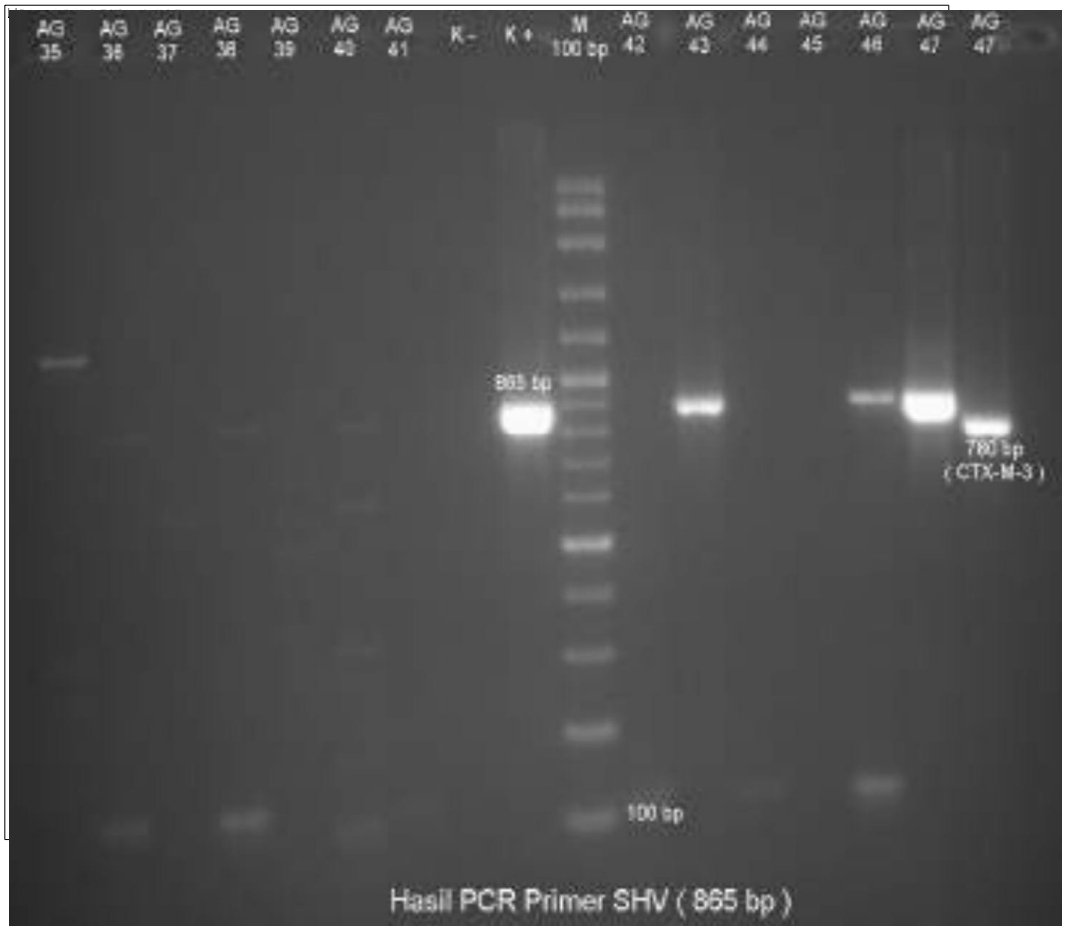
Gen SHV-12 sampel AG 1- AG17



Gen SHV-12 sampel AG 18- AG 34



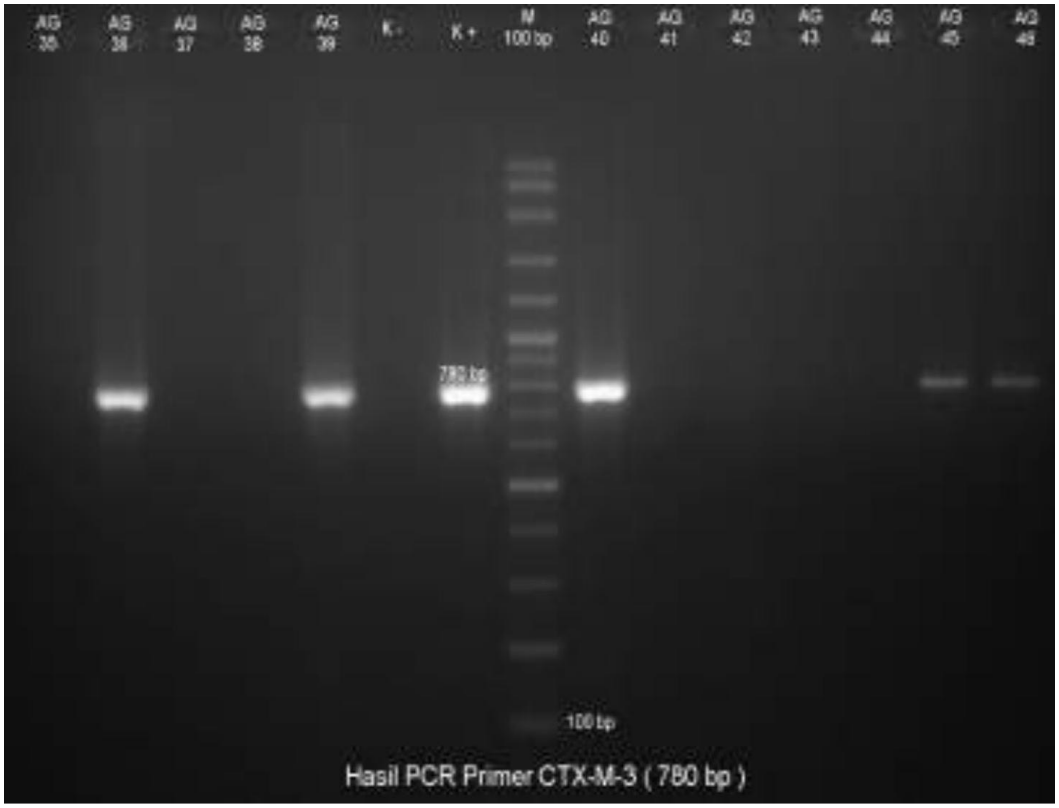
Gen SHV-12 sampel AG 35- AG 47



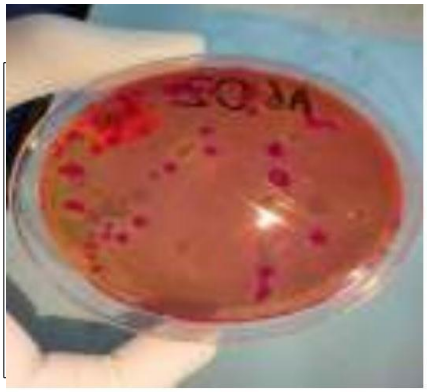
Gen CTX M-3 sampel AG 1- AG 17

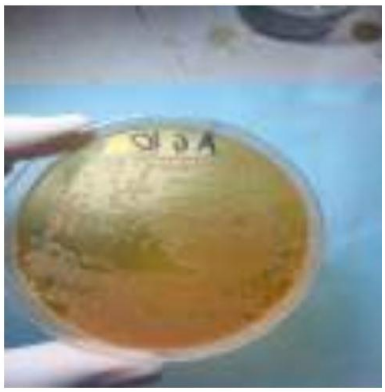
Gen CTX M-3 sampel AG 18- AG 34

Gen CTX M-3 sampel AG 35- AG 47



Lampiran 2. Dokumentasi penelitian





Koloni bakteri *Enterobacter cloacae* pada media *MacConkey* agar

Proses *running* elektroforesis

Isolat bakteri *E. cloacae* pada TSI agar (*Acid*/*acid*) dan SIM agar

Isolat bakteri *E. cloacae* pada TSI (Alkali/*acid*) dan SIM agar

Lampiran 3. Hasil identifikasi *E. cloacae* secara *fenotip* dan *genotip*

No	Kode Sampel	Vitek 2 ® *E. cloacae*	Resistansi antibiotik	Gen SHV-12	Gen CTX M-3	Kesimpulan hasil
1.	AG 01	teridentifikasi	*ampicilin*, piperacilin tazobactam, *ceftazidime*, ceftriaxone, *ampicillin sulbactam*, *ertapenem*	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12 dan CTX M-3
2.	AG 02	teridentifikasi	*ampicilin, amoxicillin clavulanat, ceftazidime, ceftriaxone, ciprofloxacin*	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12 dan CTX M-3
3.	AG 03	teridentifikasi	*cefotaxime*	positif	positif	teridentifikasi *E. cloacae*, positif gen SHV-12 dan CTX M-3
4.	AG 04	teridentifikasi	*cefotaxime, ceftazidime, ceftriaxone*	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12 dan CTX M-3
5.	AG 05	teridentifikasi	*ampicilin*, *amoxicillin clavulanat*, *cefoxitin*	negatif	positif	teridentifikasi *E. cloacae*, positif gen CTX M-3

6.	AG 06	teridentifikasi	Tidak terdapat resistansi terhadap antibiotik	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12 dan CTX M-3
7.	AG 07	teridentifikasi	*ampicilin*	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12 dan CTX M-3
8.	AG 08	teridentifikasi	tidak terdapat resistansi terhadap antibiotik	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12 dan CTX M-3
9.	AG 09	teridentifikasi	*ampicilin*, *ampicilin sulbactam*, *cefazolin*	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12 dan CTX M-3
10.	AG 10	teridentifikasi	*ampicilin*	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12 dan CTX M-3
11.	AG 11	teridentifikasi	*ampicilin*	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12 dan CTX M-3
12.	AG 12	teridentifikasi	*ceftazidime*, *ceftriaxone*	negatif	positif	teridentifikasi *E. cloacae*, positif gen CTX M-3

13.	AG 13	teridentifikasi	*amoxicillin clavulanat, levofloxacin*	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12 dan CTX M-3
14.	AG 14	teridentifikasi	*cefotaxime, ceftriaxone, gentamycin*	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12 dan CTX M-3
15.	AG 15	teridentifikasi	*ampicilin, amoxicillin clavulanate, piperacilin tazobactam, ceftazidime, ceftriaxone,*	positif	positif	teridentifikasi *E. cloacae*, positif gen SHV-12 dan CTX M-3

No	Kode Sampel	Vitek 2 ® *E.* cloacae	Resistansi antibiotik	Gen SHV-12	Gen CTX M-3	Kesimpulan hasil
16.	AG 16	teridentifikasi	tidak terdapat resistansi terhadap antibiotik	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12 dan CTX M-3
17.	AG 17	teridentifikasi	*ampicilin, cefoperazone sulbactam, meropenem*	positif	negatif	teridentifikasi *E. cloacae,* positif gen SHV-12

18.	AG 18	teridentifikasi	*ampicilin*, *amoxicillin clavulanate*	positif	negatif	teridentifikasi *E. cloacae,* positif gen SHV-12
19.	AG 19	teridentifikasi	*ceftriaxone*	positif	positif	teridentifikasi *E. cloacae*, positif gen SHV-12 dan CTX M-3
20.	AG 20	teridentifikasi	tidak terdapat resistansi terhadap antibiotik	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12 dan CTX M-3
21.	AG 21	teridentifikasi	*cefotaxime, ceftriaxone, tobramycin*	negatif	positif	teridentifikasi *E. cloacae*, positif gen CTX M-3
22.	AG 23	teridentifikasi	*ampicilin*, *ceftazidim*e, ceftriaxone, *ampicilin* sulbactam, *cefazolin, cefepime, aztreonam*	negatif	positif	teridentifikasi *E. cloacae*, positif gen CTX M-3
23.	AG 24	teridentifikasi	*ampicilin*, *ampicilin sulbactam*, *cefazolin*	Negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12 dan CTX M-3
24.	AG 25	teridentifikasi	*ampicilin*, *ceftazidim*e, ceftriaxone, *ampicilin* sulbactam, cefazolin, *cefepime*, *aztreonam*	negatif	positif	teridentifikasi *E. cloacae*, positif gen CTX M-3
25.	AG 26	teridentifikasi	*ampicilin, ceftazidime, ceftriaxone, ampicilin sulbactam, cefazolin,*	negatif	positif	teridentifikasi *E. cloacae*, positif gen CTX M-3

			cefepime, aztreonam			
26.	AG 27	teridentifikasi	*ampicilin*, piperacilin sulbactam, *ceftazidim*e, ceftriaxone, *ampicilin* sulbactam, cefazolin, Aztreonam, ertapenem,	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12 dan CTX M-3
27.	AG 28	teridentifikasi	*ampicilin*, *amoxicillin clavulanate*	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12 dan CTX M-3
28.	AG 29	teridentifikasi	*ampicilin*, *amoxicillin clavulanate*, *ampicilin* sulbactam, cefazolin	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12 dan CTX M-3
29.	AG 30	teridentifikasi	*ampicilin, ampicilin sulbactam, cefazolin*	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12dan CTX M-3

No	Kode sampel	Vitek 2 ® *E.* cloacae	Resistansi antibiotik	Gen SHV-12	Gen CTX M-3	Kesimpulan hasil
30.	AG 31	teridentifikasi	*ampicilin, piperacilin tazobactam, ceftazidime, ceftriaxone, gentamycin, ampicilin sulbactam, cefazolin, cefepime, aztreonam*	negatif	positif	teridentifikasi *E. cloacae*, positif gen CTX M-3

31.	AG 32	teridentifikasi	*ampicilin*, *ceftazidime, ceftriaxone, ampicilin sulbactam, cefazolin, aztreonam*	negatif	positif	teridentifikasi *E. cloacae*, positif gen CTX M-3
32.	AG 33	teridentifikasi	*ampicilin*, *amoxicillin clavulanate*, *piperacilin tazobactam, cefoxitin, cefotaxime, ceftazidime, ceftriaxone*	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12 dan CTX M-3
33.	AG 34	teridentifikasi	*ampicilin*, *amoxicillin clavulanate*, ceftriaxone, *gentamycin*, *ampicilin sulbactam*, *cefazolin, aztreonam*	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12 dan CTX M-3
34.	AG 35	teridentifikasi	*cefotaxime, ceftriaxone, imipenem, meropenem*	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12 dan CTX M-3
35.	AG	teridentifikasi	*ampicilin*, ceftriaxone, *gentamycin*, *ampicilin sulbactam*, cefazolin, *aztreonam*, *trimethoprim sulfamethoxazole*	negatif	positif	teridentifikasi *E. cloacae*, positif gen CTX M-3
36.	AG 37	teridentifikasi	*ampicilin*, *ampicilin sulbactam*, *cefazolin*, *ciprofloxacin*, *nitrofurantoin*	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12 dan CTX M-3
37.	AG 38	teridentifikasi	*ampicilin*, *ampicilin* sulbactam	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12dan CTX M-3
38.	AG 39	teridentifikasi	tidak terdapat resistansi terhadap antibiotik	negatif	positif	teridentifikasi *E. cloacae*, positif gen CTX M-3
39.	AG 40	teridentifikasi	*ampicilin*, *ampicilin* sulbactam, *piperacilin tazobactam*, *ceftazidime, ceftriaxone, ertapenem*	negatif	positif	teridentifikasi *E. cloacae*, positif gen CTX M-3

40.	AG 41	teridentifikasi	tidak terdapat resistansi terhadap antibiotik	positif	negatif	teridentifikasi *E. cloacae*, positif gen SHV-12
41.	AG 42	teridentifikasi	tidak terdapat resistansi terhadap antibiotik	negatif	negatif	teridentifikasi *E. cloacae*, negatif gen SHV-12 dan CTX M-3
42.	AG 43	teridentifikasi	*ampicilin*, *ampicilin* sulbactam	negatif	positif	teridentifikasi *E. cloacae*, positif gen CTX M-3
43.	AG 44	teridentifikasi	*ampicilin*, *ampicilin sulbactam*	positif	positif	teridentifikasi *E. cloacae*, positif gen SHV-12 dan CTX M-3
44.	AG 45	teridentifikasi	*ampicilin*, *ampicilin* s*ulbactam*, *cefazolin*	positif	positif	teridentifikasi *E. cloacae*, positif gen SHV-12 dan CTX M-3

Lampiran 4. Hasil identifikasi *E. cloacae* pada *MacConkey* agar, TSI agar, dan SIM agar

No.	Kode Label	Gambaran koloni pada *MacConkey* agar	Triple Sugar Iron (TSI) agar	*Sulfur Indol Motilitas (*SIM*)* agar
1.	AG 01	koloni *pink*, *mukoid*, batas tidak tegas	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
2.	AG 02	koloni besar, *pink*, *mukoid*, batas tegas	*slant= acid* *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
3.	AG 03	koloni besar, *pink*, *mukoid*	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
4.	AG 04	koloni kecil, *pink*, *mukoid*, batas tegas	*slant= acid* *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
5.	AG 05	koloni besar, *pink*, *mukoid*, batas tegas	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif

6.	AG 06	koloni kecil, *pink*, *mukoid*	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
7.	AG 07	koloni besar, *pink*, *mukoid*, batas tegas	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
8.	AG 08	koloni besar, *pink*, *mukoid*, batas tegas	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
9.	AG 09	koloni *pink*, *mukoid*, batas tidak tegas	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
10.	AG 10	koloni kecil, *pink*, *mukoid*	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
11.	AG 11	koloni *pink*, *mukoid*, batas tidak tegas	*slant= acid* *butt=acid* gas= positif	sulfur= negatif *indol*= negatif motilitas= positif

			H_2S= negatif	
12.	AG 12	koloni *pink*, *mukoid*, batas tidak tegas	*slant*= *acid* *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
13.	AG 13	koloni besar, *pink*, *mukoid*, batas tegas	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
14.	AG 14	koloni besar, *pink*, *mukoid*	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
15.	AG 15	koloni kecil, bening, *mukoid*	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
16.	AG 16	koloni *pink*, besar, *mukoid*, batas tegas	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
17.	AG 17	koloni *pink*, *mukoid*, batas tidak tegas	*slant*= alkali *butt=acid*	sulfur= negatif *indol*= negatif

			gas= positif H_2S= negatif	motilitas= positif
18.	AG 18	koloni *pink*, *mukoid*, batas tidak tegas	*slant*= *acid* *butt*=*acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
19.	AG 19	koloni *pink*, *mukoid*, batas tidak tegas	*slant*= alkali *butt*=*acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
20.	AG 20	koloni *pink*, besar *mukoid*, batas tegas	*slant*= alkali *butt*=*acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
21.	AG 21	koloni *pink*, *mukoid*, batas tegas	*slant*= alkali *butt*=*acid* gas= positif H_2S= negatif	gulfur= negatif *indol*= negatif motilitas= positif
22.	AG 23	koloni *pink*, *mukoid*, besar	*slant*= alkali *butt*=*acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif

23.	AG 24	koloni *pink*, *mukoid*, batas tidak tegas	*slant*= *acid* *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
24.	AG 25	koloni *pink*, *mukoid*, batas tidak tegas	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
25.	AG 26	koloni *pink*, *mukoid*, batas tidak tegas	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
26.	AG 27	koloni *pink*, *mukoid*, batas tidak tegas	*slant*= *acid* *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
27.	AG 28	koloni *pink*, *mukoid*, batas tidak tegas	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
28.	AG 29	koloni *pink*, *mukoid*, batas tidak tegas	*slant*= *acid* *butt=acid* gas= positif	sulfur= negatif *indol*= negatif motilitas= positif

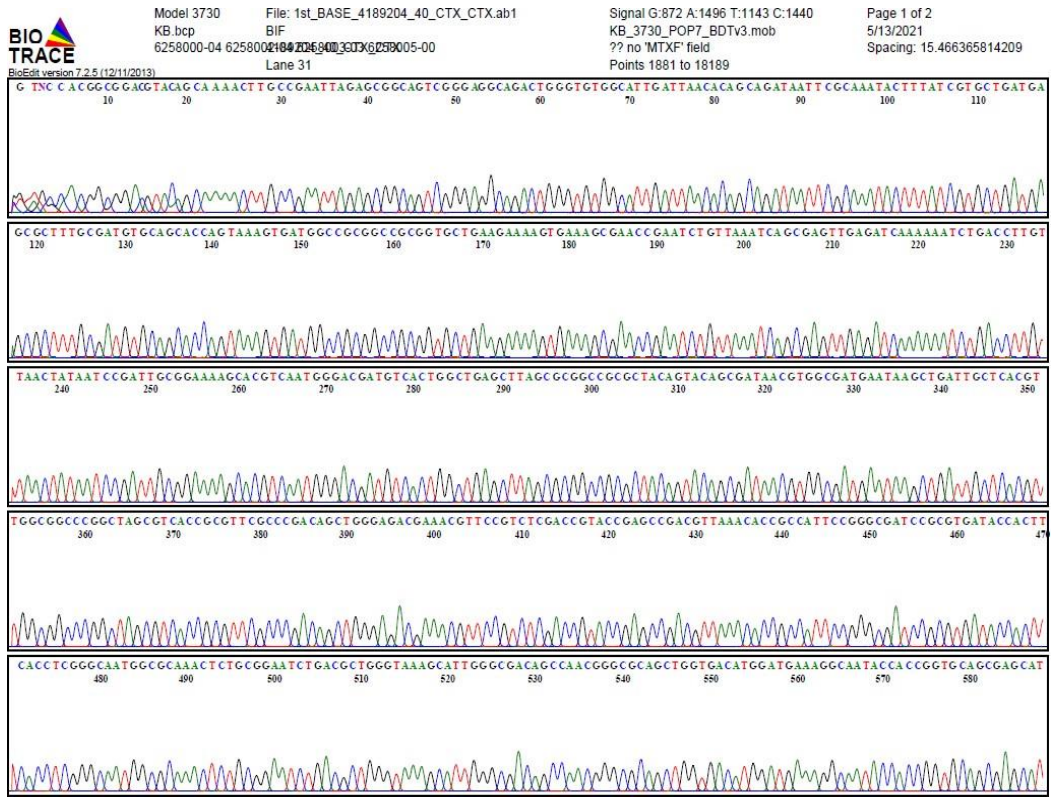
			H_2S= negatif	
29.	AG 30	koloni *pink*, *mukoid*, batas tidak tegas	*slant*= *acid* *butt*=*acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
30.	AG 31	koloni *pink*, *mukoid*, batas tidak tegas	*slant*= alkali *butt*=*acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
31.	AG 32	koloni *mukoid*, kecil, batas tidak tegas	*slant*= alkali *butt*=*acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
32.	AG 33	koloni *mukoid*, *pink*, koloni besar	*slant*= *acid* *butt*=*acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
33.	AG 34	koloni *pink*, *mukoid*, batas tidak tegas	*slant*= *acid* *butt*=*acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
34.	AG 35	koloni *pink*, *mukoid*, batas tidak tegas	*slant*= *acid* *butt*=*acid*	sulfur= negatif *indol*= negatif

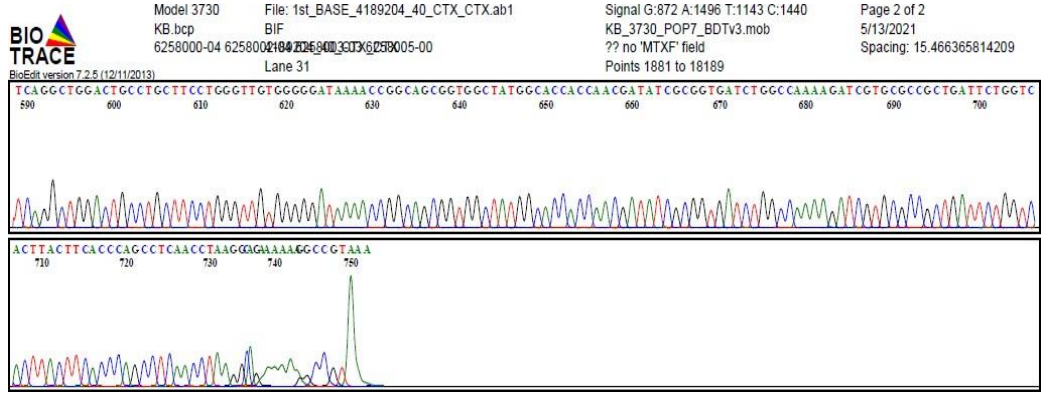
			gas= positif H_2S= negatif	motilitas= positif
35.	AG 36	koloni *pink*, *mukoid*, batas tidak tegas	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
36.	AG 37	koloni *pink*, *mukoid*, batas tidak tegas	*slant*= *acid* *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
37.	AG 38	koloni *pink*, *mukoid*, batas tidak tegas	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
38.	AG 39	koloni *pink*, *mukoid*, koloni besar, batas tidak tegas	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
39.	AG 40	koloni *pink*, *mukoid*, batas tidak tegas	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif

40.	AG 41	koloni *pink*, *mukoid*, batas tidak tegas	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
41.	AG 42	koloni *mukoid*, bening, kecil	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
42.	AG 43	koloni *mukoid*, kecil	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
43.	AG 44	koloni *pink*, *mukoid*, besar	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif
44.	AG 45	koloni *pink*, *mukoid*, besar, batas tegas	*slant*= alkali *butt=acid* gas= positif H_2S= negatif	sulfur= negatif *indol*= negatif motilitas= positif

Lampiran 5. Hasil *sequensing* gen CTX M-3 dan gen SHV-12

5.1 *sequensing* gen CTX M-3





gen CTX M-3

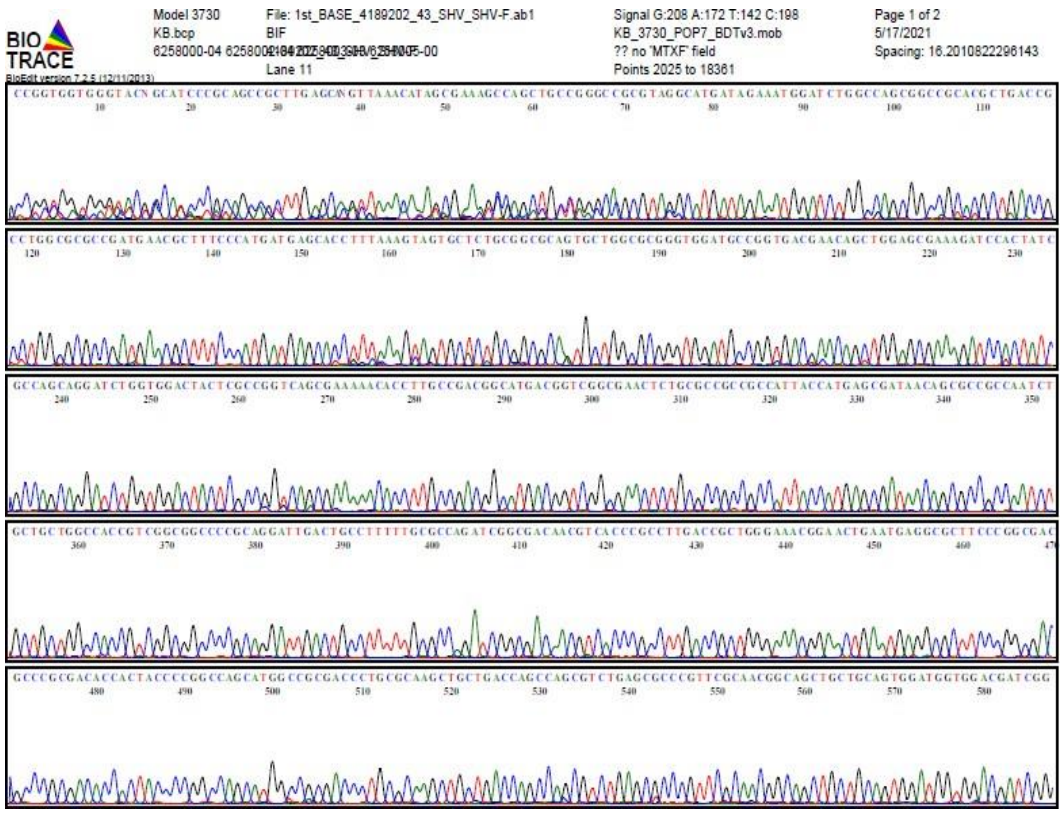
Sequences producing significant alignments:

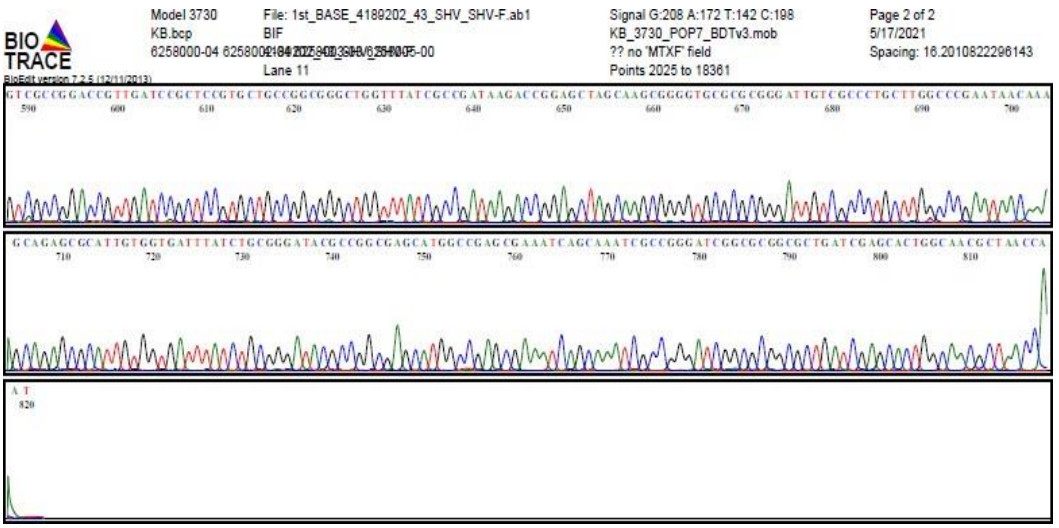
Select for downloading or viewing reports	Description	Scientific Name	Max Score	Total Score	Query Cover	E value	Per. Ident	Acc. Len	Accession
Select seq MK736669.1	Enterobacter cloacae strain EclC2185 plasmid pEclC2185CTXM15, complete sequence	Enterobacter cloacae	1354	1354	97%	0.0	99.86%	305912	MK736669.1
Select seq MK559056.1	Enterobacter cloacae strain CH8(2) plasmid CTX-M family beta-lactamase (blaCTX-M) gene, partial cds	Enterobacter cloacae	1354	1354	97%	0.0	99.73%	873	MK559056.1
Select seq CP040827.1	Enterobacter cloacae strain NH77 chromosome, complete genome	Enterobacter cloacae	1354	1354	97%	0.0	99.86%	5040532	CP040827.1
Select seq MG844171.1	Enterobacter cloacae strain ESP110 plasmid insertion sequence ISEcp1 ISEcp1 transposase (tnpA) gene, complete cds; and class A extended-spectrum beta-lactamase CTX-M-15 (blaCTX-M) and tryptophan synthase subunit beta-like protein genes, complete cds	Enterobacter cloacae	1354	1354	97%	0.0	99.73%	2959	MG844171.1
Select seq KY346922.1	Enterobacter cloacae strain ZZEN540 CTX-M-15 beta-lactamase gene, complete cds	Enterobacter cloacae	1354	1354	97%	0.0	99.73%	876	KY346922.1
Select seq KY346921.1	Enterobacter cloacae strain ZZEN513 CTX-M-15 beta-lactamase gene, complete cds	Enterobacter cloacae	1354	1354	97%	0.0	99.73%	876	KY346921.1

Sequences producing significant alignments:

Select for downloading or viewing reports	Description	Scientific Name	Max Score	Total Score	Query Cover	E value	Per. Ident	Acc. Len	Accession
Select seq KY346934.1	Enterobacter cloacae strain ZZEN521 CTX-M-3 beta-lactamase gene, complete cds	Enterobacter cloacae	1349	1349	97%	0.0	99.59%	876	KY346934.1
Select seq MF062700.1	Enterobacter cloacae strain T5282 plasmid pT5282-CTXM, complete sequence	Enterobacter cloacae	1349	1349	97%	0.0	99.73%	60206	MF062700.1
Select seq KX015668.1	Enterobacter cloacae strain CY01 plasmid pCY-CTX, complete sequence	Enterobacter cloacae	1349	1349	97%	0.0	99.73%	116700	KX015668.1
Select seq AP022128.1	Enterobacter cloacae plasmid pWP5-S18-CRE-02_2 DNA, complete genome, strain: WP5-S18-CRE-02	Enterobacter cloacae	1349	1349	97%	0.0	99.59%	92491	AP022128.1
Select seq HQ214050.1	Enterobacter cloacae strain S-595 plasmid insertion sequence ISEcp1, partial sequence; and beta-lactamase CTX-M-3 (blaCTX-M-3) and hypothetical protein genes, complete cds	Enterobacter cloacae	1349	1349	97%	0.0	99.59%	1498	HQ214050.1
Select seq HQ214046.1	Enterobacter cloacae strain S-361 plasmid IncF::L/M insertion sequence ISEcp1 transposase (tnpA) gene, complete cds; and beta-lactamase CTX-M-3 (blaCTX-M-3) and hypothetical protein genes, complete cds	Enterobacter cloacae	1349	1349	97%	0.0	99.59%	3022	HQ214046.1
Select seq AY954529.1	Enterobacter cloacae plasmid pEC002 beta-lactamase CTX-M-22 (blaCTX-M-22) gene, complete cds	Enterobacter cloacae	1349	1349	97%	0.0	99.59%	876	AY954529.1

5.2 *sequensing* gen SHV-12





gen SHV-12

Sequences producing significant alignments:

Select for downloading or viewing reports	Description	Scientific Name	Max Score	Total Score	Query Cover	E value	Per. Ident	Acc. Len	Accession
Select seq MH399264.1	Enterobacter cloacae strain RJ702 plasmid pIMP26, complete sequence	Enterobacter cloacae	1450	1450	97%	0.0	99.50%	329420	MH399264.1
Select seq MH460797.1	Enterobacter cloacae strain KMB37 extended spectrum beta-lactamase SHV-12 (blaSHV) gene, blaSHV-12 allele, complete cds	Enterobacter cloacae	1450	1450	97%	0.0	99.37%	861	MH460797.1
Select seq MH460793.1	Enterobacter cloacae strain KSR2 extended spectrum beta-lactamase SHV-12 (blaSHV) gene, blaSHV-12 allele, complete cds	Enterobacter cloacae	1450	1450	97%	0.0	99.37%	861	MH460793.1
Select seq MF415608.1	Enterobacter cloacae strain hhy03 plasmid pNDM-BJ03, complete sequence	Enterobacter cloacae	1450	1450	97%	0.0	99.50%	60125	MF415608.1
Select seq MF402906.1	Enterobacter cloacae strain ZUECL4 extended-spectrum beta-lactamase SHV-12 (blaSHV-12) gene, complete cds	Enterobacter cloacae	1450	1450	97%	0.0	99.37%	861	MF402906.1
Select seq MG653177.1	Enterobacter cloacae strain BVT29 extended spectrum beta lactamase (SHV) gene, SHV-12 allele, complete cds	Enterobacter cloacae	1450	1450	97%	0.0	99.37%	861	MG653177.1
Select seq MG653176.1	Enterobacter cloacae strain BVT22 extended spectrum beta lactamase (SHV) gene, SHV-12 allele, complete cds	Enterobacter cloacae	1450	1450	97%	0.0	99.37%	861	MG653176.1

Sequences producing significant alignments:

Select for downloading or viewing reports	Description	Scientific Name	Max Score	Total Score	Query Cover	E value	Per. Ident	Acc. Len	Accession
Select seq MF344583.1	Enterobacter cloacae strain N1863 plasmid pN1863-HI2, complete sequence	Enterobacter cloacae	1450	1450	97%	0.0	99.50%	349834	MF344583.1
Select seq KY270852.1	Enterobacter cloacae strain T5282 plasmid pT5282-mphA, complete sequence	Enterobacter cloacae	1450	1450	97%	0.0	99.37%	282423	KY270852.1
Select seq KY296103.1	Enterobacter cloacae strain 13E169 plasmid pHN84NDM, complete sequence	Enterobacter cloacae	1450	1450	97%	0.0	99.37%	53771	KY296103.1
Select seq NG_050590.1	Enterobacter cloacae CRE727 pNDM-HF727 blaSHV gene for class A extended-spectrum beta-lactamase SHV-12, complete CDS	Enterobacter cloacae	1450	1450	97%	0.0	99.37%	1061	NG_050590.1
Select seq KP975077.1	Enterobacter cloacae strain MRSN17626 plasmid pMRVIM0813, complete sequence	Enterobacter cloacae	1450	2894	97%	0.0	99.50%	311662	KP975077.1
Select seq KF976405.1	Enterobacter cloacae strain CRE727 plasmid pNDM-HF727, complete sequence	Enterobacter cloacae	1450	1450	97%	0.0	99.37%	54035	KF976405.1
Select seq AP022520.1	Enterobacter cloacae STN0717-73 plasmid pSTN0717-73-1 DNA, complete sequence	Enterobacter cloacae	1450	1450	97%	0.0	99.37%	266476	AP022520.1
Select seq	Enterobacter cloacae strain B02/0225 beta-lactamase (blaSHV) gene, partial cds	Enterobacter cloacae	1450	1450	97%	0.0	99.37%	860	JN037849.1

KEMENTERIAN PENDIDIKAN DAN KEBUDAYAAN
UNIVEkSITAS HASANUDDIN FAKULTAS KEDOKTERAN
KOMITE ETIK PENELITIAN KESEHATAN
RSPTN UNtVERSITAS HASANUDDtN
RSUP Dr. \X/AHIDIN SUDIROHUSODO MAKASSAR
Sekretariat : Lantai 2 Gedung Laboratorium Terpadu
JL.PERINTIS KEMERDEKAAN KAMPUS TAMALANREA KM.IO MAKASSAR 90245.
Contact Person: dr. Agussalim Bukhari.,MMed,PhD. SpGK TELP. 081241850858, 04115780103, Fax : 0411-581431

REKOMENDASI PERSETUJUAN ETIK

Nomor : 149/UN4.6.4.5.31/ PP36/ 2021

Tanggal: 8 Maret 2021

Dengan ini Menyatakan bahwa Protokol dan Dokumen yang Berhubungan Dengan Protokol berikut ini telah mendapatkan Persetujuan Etik :

No Protokol	UH21020124	No Sponsor Protokol	
Peneliti Utama	dr. Donnaries Hangga Kusuma	Sponsor	
Judul Peneliti	Distribusi gen CTX M-3 & SHV-12 pada isolate Enterobactercloacae di RSUH dan RSWS Makassar		
No Versi Protokol	1	Tanggal Versi	2 Maret 2021
No Versi PSP		Tanggal Versi	
Tempat Penelitian	RS Universitas Hasanuddin dan RS Dr. Wahidin Sudirohusodo Makassar		
Jenis Review	[X] Exempted [ ] Expedited [ ] Fullboard Tanggal	Masa Berlaku 8 Maret 2021 sampai 8 Maret 2022	Frekuensi review lanjutan
Ketua Komisi Etik Penelitian Kesehatan FKUH	Nama Prof.Dr.dr. Suryani As'ad, M.Sc.,Sp.GK (K)	Tanda tangan	
Sekretaris Komisi Etik Penelitian Kesehatan FKUH	Nama dr. Agussalim Bukhari, M.Med.,Ph.D.,Sp.GK I)	Tanda tangan	

Kewajiban Peneliti Utama:

- Menyerahkan Amandemen Protokol untuk persetujuan sebelum di implementasikan
- Menyerahkan Laporan SAE ke Komisi Etik dalam 24 Jam dan dilengkapi dalam 7 hari dan Lapor SUSAR dalam 72 Jam setelah Peneliti Utama mv•nerirna laporan
- Menyerahkan Laporan Kemajuan (progress report) setiap 6 bulan untuk penelitian resiko tinggi dan setiap setahun untuk penelitian resiko rendah
- Menyei ahkan laporan akhir setelah Penelitian berakhir
- Melaporkan penyimpangan dari prokol yang disetujui (protocol deviation / violation)
- Mematuhi semua peraturan yang ditentukan "