SUPPLEMENTARY DATA

Supplementary Table 1: Example search strategy

Search	Query
#4	Search (#1 AND #2 AND #3)
#3	Search "HIV Infections" [MeSH] OR HIV [MeSH] OR HIV [tiab] OR hiv-1*[tiab]
	OR hiv1[tiab] OR HIV infect*[tiab] OR human immunodeficiency
	virus[tiab] OR human immunedeficiency virus[tiab] OR human immuno-deficiency
	virus[tiab] OR human immunedeficiency virus[tiab] OR ((human immun*) AND
	(deficiency virus[tiab])) OR acquired immunodeficiency syndrome[tiab] OR acquired
	immunedeficiency syndrome[tiab] OR acquired immuno-deficiency syndrome[tiab] OR
	acquired immune-deficiency syndrome[tiab] OR ((acquired immun*) AND (deficiency
	syndrome[tiab]))
#2	Search "second line"[tiab] OR "second-line"[tiab] or "second line treatment"[tiab] OR
	"second-line treatment" [tiab] or "second-line antiretroviral therapy" [tiab] OR "second
	line antiretroviral therapy"[tiab] OR "protease inhibitor*"[tiab] OR "HIV Protease
	Inhibitor" [MeSH] OR "integrase inhibitor" [tiab] OR "HIV Integrase Inhibitors" [MeSH]
	OR lopinavir[tiab] OR lopinavir/ritonavir[tiab] OR atazanavir[tiab] OR darunavir[tiab]
	OR ritonavir[tiab] OR raltegravir[tiab] OR dolutegravir[tiab] OR indinavir[tiab] OR
	nelfinavir[tiab] OR fosamprenavir[tiab] OR etravirine[tiab] OR "treatment
	failure"[MeSH] OR stribild[tiab] OR eviplera[tiab] OR complera[tiab] OR
11.1	maraviroc[tiab]
#1	Search Africa[MeSH] OR africa[tiab] OR "sub Saharan africa"[tiab] OR "sub-Saharan
	Africa"[tiab] OR Angola[tiab] OR Benin[tiab] OR Botswana[tiab] OR "Burkina
	Faso"[tiab] OR Burundi[tiab] OR Cameroon[tiab] OR "Cape Verde"[tiab] OR
	"Central African Republic"[tiab] OR Chad[tiab] OR Comoros[tiab] OR "Republic of the Congo"[tiab] OR "Democratic Republic of the Congo"[tiab] OR "Cote
	d'Ivoire"[tiab] OR Djibouti[tiab] OR "Equatorial Guinea"[tiab] OR Eritrea[tiab] OR
	Ethiopia[tiab] OR Gabon[tiab] OR "The Gambia"[tiab] OR Ghana[tiab] OR
	Guinea[tiab] OR "Guinea-Bissau"[tiab] OR Kenya[tiab] OR Lesotho[tiab] OR
	Liberia[tiab] OR Madagascar[tiab] OR Malawi[tiab] OR Mali[tiab] OR
	Mauritania[tiab] OR Mauritius[tiab] OR Mozambique[tiab] OR Namibia[tiab] OR
	Niger[tiab] OR Nigeria[tiab] OR Rwanda[tiab] OR "Sao Tome and Principe"[tiab] OR
	Senegal[tiab] OR Seychelles[tiab] OR "Sierra Leone"[tiab] OR "South Africa"[tiab]
	OR "South Sudan"[tiab] OR Sudan[tiab] OR Swaziland[tiab] OR Tanzania[tiab] OR
	Togo[tiab] OR Uganda[tiab] OR Zambia[tiab] OR Zimbabwe[tiab]
	1 1050[thto] Or Oganica[thto] Or Zamola[thto]

[tiab] = title and abstract [MeSH]= medline search heading

Supplementary Table 2: Outcomes of second-line ART at 48 weeks, ITT analysis

Study/Reference	Number of	Virological ou	itcome, n (%)	No virolo		Alive, receiving			
•	participants	Suppression	LLV	VF	Dead	TO	LTFU	Missing data	IFU	care, n (%)
Randomised controlled t	rials									
La Rosa[1]	162	148 (91.4)	2 (1.2)	4 (2.5)	2 (1.2)	0(0)	3 (1.9)	3 (1.9)	_	157/162 (96.9)
Ciaffi[2]	451	399 (88.5)	11 (2.4)	29 (6.4)	6 (1.3)	0(0)	4 (0.9)	2 (0.4)	-	441/451 (97.8)
Paton[3]	426	336 (78.9)	14 (3.3)	45 (10.6)	21 (4.9)	0(0)	0 (0)	10 (2.3)	-	405/426 (95.1)
Boyd[4]/Amin[5]	100	84 (84.0)	1 (1.0)	8 (8.0)	4 (4.0)	1 (1.0)	1 (1.0)	1 (1.0)	-	94/100 (94.0)
Gross[6]	132	112 (84.8)	3 (2.2)	9 (6.8)	4 (3.0)	1 (0.8)	0 (0)	3 (2.3)	-	127/132 (96.2)
Prospective/ retrospectiv	e observation	al cohort								
Osinusi-	73	36 (49.3)	-	3 (4.1)	3 (4.1)	5 (6.8)	4 (5.5)	22 (30.1)	0 (0)	61/73 (83.6)
Adekanmbi[7]*										
Shearer[8]*	1150	694 (60.3)	-	233(20.2)	22 (1.9)	0(0)	123 (10.9)	223 (19.3)	0 (0)	1005/1150 (87.4)
Schoffelen[9]	156	68 (43.6)	2 (1.3)	28 (17.9)	4 (2.6)	4 (2.6)	17 (10.9)	30 (21.2)	3 (1.9)	128/153 (83.7)
Wandeler[10]	971	253 (26.1)	21 (2.2)	120 (12.4)	22 (2.3)	0(0)	49 (5.0)	295 (30.3)	211 (21.7)	689/760 (90.7)
Boender[11]/Sigaloff[1	243	178 (73.3)	8 (3.3)	20 (8.2)	10 (4.1)	5 (2.1)	15 (6.1)	7 (2.8)	0 (0)	213/243 (87.7)
2]										
Murphy[13]	136	94 (69.1)	0 (0)	26 (19.1)	0(0)	6 (4.4)	4 (2.9)	6 (4.4)	0 (0)	126/136 (92.6)
Johnston[14, 15]*	417	174 (41.7)	-	111 (26.6)	19 (4.6)	23 (5.6)	29 (7.0)	61 (14.6)	-	346/417 (83.0)
Hosseinipour[16]	101	75 (74.2)	-	13 (12.8)	10 (9.9)	0 (0)	3 (3.0)	0 (0)	0(0)	88/101 (87.1)
Castelnuovo[17]	40	30 (75.0)	0 (0)	9 (22.5)	1 (2.5)	0 (0)	0 (0)	0 (0)	0 (0)	39/40 (97.5)

ITT= Intention-to-treat; LLV= low level viraemia (VL 400-999 copies/ml); VF= Virological failure (>1000 copies/ml); TO= transferred out; LTFU= Lost to follow up; IFU= Insufficient follow up the proportion of patients in prospective observational cohorts who had not accrued sufficient follow-up at the time of analysis.

^{*}In the indicated cohorts, viral load data was dichotomised at the 400 copies/ml cut-off; data was not available for the proportion with LLV; VF is defined as >400 copies/ml

Supplementary Table 3: Outcomes of second-line ART at 96 weeks, ITT analysis

Study/ Reference	Number of	Virolog	ical outcome	e, n (%)	No virolo	ogical data,		Alive, receiving		
	participants	Suppression	LLV	VF	Dead	TO	LTFU	Missing data	IFU	care
Randomised controlled t	trials									
Paton[3]	426	326 (76.5)	7 (1.6)	46 (10.8)	30 (7.0)	0 (0)	4 (0.9)	13 (3.1)	-	392/426 (92.0)
Amin[5]	100	76 (76.0)	5 (5.0)	9 (9.0)	5 (5.0)	1 (1.0)	2 (2.0)	2 (2.0)	-	92/100 (92.0)
Prospective/ retrospective	ve observation	nal cohort								
Osinusi-Adekanmbi[7]	73	41 (56.2)	0(0)	3 (4.1)	5 (6.8)	6 (8.2)	6 (8.2)	12 (16.4)	0 (0)	56/73 (76.7)
Schoffelen[9]	156	49 (31.4)	2 (1.3)	15 (9.6)	8 (5.1)	9 (5.8)	24 (15.4)	10 (6.4)	39 (25.0)	76/117 (65.0)
Wandeler[10]	971	152 (15.7)	7 (0.7)	58 (6.0)	39 (4.0)	0 (0)	67 (6.9)	204 (21.0)	444 (45.7)	421/527 (80.0)
Boender[11]/Sigaloff[1	243	150 (61.7)	0 (0)	27 (11.1)	14 (5.8)	13 (5.3)	24 (9.9)	15 (6.2)	0 (0)	192/243 (79.0)
2]										
Murphy[13]	136	74 (54.4)	0(0)	6 (4.4)	1 (0.7)	19 (14.0)	17 (12.5)	19 (14.0)	0 (0)	99/136 (72.8)
Castelnuovo[17]	40	34 (85.0)	0(0)	3 (7.5)	1 (2.5)	0(0)	2 (5.0)	0 (0)	0 (0)	37/40 (92.5)

ITT= Intention-to-treat; LLV= low level viraemia (VL 400-999 copies/ml); VF= Virological failure (>1000 copies/ml); TO= transferred out; LTFU= Lost to follow up; IFU= Insufficient follow up the proportion of patients in prospective observational cohorts who had not accrued sufficient follow-up at the time of analysis.

Supplementary Table 4: Quality assessment of included studies

Study/ Ref	Participant characteristic s adequately described?	Eligibility criteria explicit & appropriate?	Participant recruitment/ selection adequately described?	Intervention clearly described?	Co- intervention s clearly reported?	Outcome measure defined?	Outcomes measured before & after intervention ?	Length of follow up reported?	Loss to follow up reported?	Adverse events reported?	Competing interests/ support reported?
La Rosa[1]	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y
Ciaffi[2]	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y
Paton[3]	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y
Boyd[4]/ Amin[5]	Y	Y	Y	Y	N	Y	Y	Y	Y	Y	Y
Gross[6]	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y
Osinusi- Adekanmbi[7]	Partially reported	Y	Y	Y	N	Y	Y	Y	Y	N	Y
Shearer[10]	Partially reported	Y	Y	Y	Y	Y	Y	Y	Y	N	Y
Schoffelen[9]	Y	Y	Y	Y	N	Y	Y	Y	Y	N	Y
Adetunji[18]a	Partially reported	Y	Y	N	N	Y	Y	Y	N^a	N	Y
Wandeler[10]	Partially reported	Y	Y	N	N	Y	Y	Y	Y	N	Y
Boender[11]/ Sigaloff[12]	Y	Y	Y	N	N	Y	Y	Y	Y	N	Y
Murphy[13]	Partially reported	Y	Y	N	N	Y	Y	Y	Y	N	Y
Johnston[14, 15]	Y	Y	Y	N	Y	Y	Y	Y	Y	Y	Partially reported
Hosseinipour[16]	Y	Y	Y	Y	N	Y	Y	Y	Y	Y	Partially reported
Castelnuovo[1 7]	Partially reported	N	N	Y	N	Y	Y	Y	Y	Y	Y

Abbreviations: Y yes; N no.

^aAdetunji et al included only participants with a known virological outcome at 12months; therefore study data was included only in on-treatment analyses

Supplementary Table 5: On-treatment outcomes of second-line ART at 48 and 96 weeks

Study/ Reference	Virolog	gical outcomes at 48	weeks, n (%)		Virological outcomes at 96 weeks, n (%)				
•	Total	Suppression	LLV	VF	Total	Suppression	LLV	VF	
Randomised controlled tri	als								
La Rosa[1]	154	148 (96.1)	2 (1.3)	4 (2.5)					
Ciaffi[2]	439	399 (90.9)	11 (2.5)	29 (6.6)					
Paton[3]	395	336 (85.1)	14 (3.5)	45 (11.4)	379	326 (86.0)	7 (1.8)	46 (12.1)	
Boyd[4]/ Amin[5]	93	84 (90.3)	1 (1.1)	8 (8.6)	90	76 (84.4)	5 (5.6)	9 (10.0)	
Gross[6]	124	112 (90.3)	3 (2.4)	9 (7.3)					
Prospective/ retrospective	observati	onal cohort							
Osinusi-Adekanmbi[7]*	39	36 (92.3)	-	3 (7.7)	44	41 (93.2)	-	3 (6.8)	
Shearer[8]*	927	694 (74.9)	-	233 (25.1)					
Schoffelen[9]	98	68 (69.4)	2 (2.0)	28 (28.6)	66	49 (74.2)	2 (3.0)	15 (22.7)	
Wandeler[10]	394	253 (64.2)	21 (5.3)	120 (30.5)	217	152 (70.0)	7 (3.2)	58 (26.7)	
Adetunji[18] ^a	225	191 (84.9)	0 (0)	34 (15.1)					
Boender[11]/	206	178 (86.4)	8 (3.9)	20 (9.7)	177	150 (84.7)	0 (0)	27 (15.3)	
Sigaloff[12]									
Murphy[13]	120	94 (78.3)	0 (0)	26 (21.7)	80	74 (92.5)	0 (0)	6 (7.5)	
Johnston[14, 15]*	285	174 (61.1)	-	111 (38.9)					
Hosseinipour[16]*	88	75 (85.2)	-	13 (14.8)					
Castelnuovo[17]	39	30 (76.9)	0 (0)	9 (23.1)	37	34 (91.9)	0 (0)	3 (8.1)	

LLV low level viraemia (VL 400-1000 copies/ml); VF= Virological failure;

^{*}In the indicated cohorts, viral load data was dichotomised at 400 copies/ml; data were not available for the proportion with LLV; VF is defined as >400 a In Adetjunji *et al*, only patients with a known viral load at 12 months were included; the study is therefore only included in the on-treatment analysis.

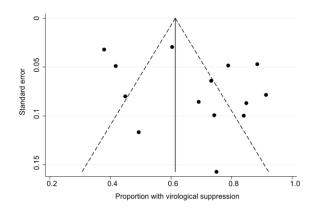
Supplementary Table 6: List of baseline NRTI mutations in included studies

Study	Duration 1st line ART (months), median (IQR)	Sequences available	GenBank accession #a	M184VI, n(%)	TAMs, n(%)
Paton[19]	48 (34,65)	787ª	KY061369 KY062155	730 (92.7)	585 (74.3)
Sigaloff[12]	27 (15, 44)	183	JN132214- JN132396	150 (82.0)	92 (50.3)
Boyd[20]	40 (22, 65)	91	-	77 (84.6)	25 (27.47)
Castelnouvo[17]	22 (19, 29)	16	-	13 (81.3)	2 (12.5)
Hosseinipour[16, 21]	37 (27, 50)	96	-	77 (80.2)	54 (56.3)
Johnston[14]	18 (10, 27)	115	KC921018- KC921144	77 (67.0)	39 (33.9)

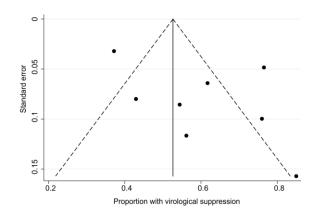
^a Sequences available at: https://www.ncbi.nlm.nih.gov/nucleotide/ Abbreviations: NRTI nucleoside reverse transcriptase inhibitor, IQR Interquartile range, TAM Thymidine analogue mutations, according to the Stanford HIV database (8.2)

Supplementary Figure 1: Funnel plots of studies included in intention to treat analysis of virologic suppression at 48 weeks (A) and 96 weeks (B)

A: 48 weeks



B: 96 weeks



Supplementary Material: Quality assessment checklist

Adapted from: Development of a quality appraisal tool for case series using a modified Delphi technique (Institute of Health Economics) 2012[22]

Study population

1. Are the characteristics of the participants included in the study described?

Yes: The authors should report the total number, age, and gender distribution of the participants commencing second line ART, the duration and type of prior ART use, CD4 count and viral load at start of 2^{nd} line therapy, if these criteria were used to define treatment failure.

Partially reported: The criteria above are incompletely reported

No: None of the relevant characteristics of the participants is reported.

2. Are the eligibility criteria (inclusion and exclusion criteria) to entry the study explicit and appropriate?

Yes: The eligibility criteria are clearly stated and replicable. A statement on age of eligibility, type of prior ART use accepted, definition of 1st line treatment failure are required.

No: The eligibility criteria are not clearly stated or are inappropriate.

3. Was the method of recruitment adequately described and appropriate?

Yes: There is a clear statement of the method of selection of participants. If viral load testing occurred, all patients received viral load testing or a clear description of the reason for missing data was given. For resistance studies, all participants who were virologically failing underwent resistance genotyping or they were a random unbiased selection of those failing and consistent virological thresholds for sequencing were applied.

Unclear: The method used to recruit participants is not clearly stated or no information is provided about the method used to recruit participants in the study. The method of selection for viral load testing was not described. Selection for resistance genotyping was vague or uncertain.

No: The participants were recruited based on other criteria, such as access to intervention or availability of resources, that could lead to a selection bias. Viral load or genotyping was not random and depended on availability or access to resources.

Intervention and co-intervention

4. Was the intervention clearly described in the study?

Yes: There is a detailed description about the characteristics of the intervention. The type and where non-standard, the dose and frequency of 2nd line ART should be described including details on choice of NRTIs

No: The information provided is unclear, or important parameters of the intervention are missing from the presentation.

5. Were additional interventions (co-interventions) adequately reported in the study?

Yes: Co-interventions such as co-trimoxazole preventative therapy and any additional adherence support or increased clinic monitoring provided alongside 2nd line therapy should be described or if routine care was offered this should be described.

No: The use of co-interventions is not adequately described.

Outcome measures

6. Are the outcome measures clearly defined in the introduction or methods section?

Yes: All relevant (primary and secondary) outcomes that match the objective(s) of the study are described in the introduction or methods section. If virological/clinical/immunological failure, this must be defined (threshold, type and timing of tests). If loss to follow up is provided as an outcome this should be defined.

Partially reported: Some of the relevant outcomes are briefly reported in the introduction or methods section.

No: The outcomes are reported for the first time in the results or conclusion section of the study. The relevant outcomes are briefly mentioned without any details in the results, discussion, or conclusion section(s). The outcomes reported are not relevant to study objective(s).

7. Were outcomes measured before and after intervention?

Yes: The relevant outcomes (viral load) are measured before and after applying the intervention.

Unclear: It is unclear when the outcomes were measured.

No: The outcomes are measured only after applying the intervention.

Results and conclusions

8. Was the length of follow-up reported?

Yes: The length of follow-up is clearly reported.

Unclear: The duration of follow-up is not clearly reported.

No: The length of follow-up is not reported, or the duration of the study is unclear.

9. Was the loss to follow-up reported?

Yes: The number or proportion of patients lost to follow-up is reported.

Unclear: It is not clear from the information provided how many participants were lost to follow-up or it is an inconsistence of reporting lost to follow-up (e.g. discrepancies between information from tables and text).

No: The number or proportion of patients lost to follow-up is not reported.

10. Are adverse events reported?

Yes: The undesirable or unwanted consequences of the intervention during the study period or within a prespecified time period are reported. Absence of any adverse event(s) is acknowledged in the study.

Partially reported: It is deducible that only some but not all potential adverse events are reported. **No**: There is no statement about the presence or absence of adverse events.

Competing interest and source of support

11. Are both competing interest and source of support for the study reported?

Yes: Both competing interest and source of support (financial or other) received for the study are reported, or the absence of any competing interest and source of support is acknowledged.

Partially reported: Only one of these elements is reported.

No: Either there is no information available about competing interests and sources of support, or only one of these elements is reported.

Supplementary Material: List of major protease resistance mutations

The following mutations were designated as major protease resistance mutations in this study, as defined by the Stanford HIV database[23]:

D30N, V32I, L33F, M46IL, I47VA, G48VM, I50LV, I54VTALM, L76V, V82ATFS, I84V, N88S and L90M

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