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Moving towards a person-centred HIV care cascade: An exploration of potential biases and errors in routine data in South Africa

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Abstract

In 2022, in recognition of lags in data infrastructure, the World Health Organisation (WHO) recommended the use of routinely linked individual patient data to monitor HIV programmes. The WHO also recommended a move to person-centred care to better reflect the experiences of people living with HIV. The switch from aggregated service level data to person-centred data will likely introduce some biases and errors. However, little is understood about the direction and magnitude of these biases. We investigated HIV-testing and HIV-care engagement from 2014 to 2018 in the Agincourt Health and Demographic Surveillance System (HDSS). We digitised and linked HIV patient clinic records to HDSS population data in order to estimate biases in routine clinical data. Using this linked data, we followed all individuals linked to HIV-related clinic data throughout their care pathway. We built sequences to represent these pathways. We performed sequence and cluster analyses for all individuals to categorise patterns of care engagement and identified factors associated with different engagement patterns using multinomial logistic regression. Our analyses included 4947 individuals who were linked to 5084 different patient records. We found that routine data would have inflated patient numbers by 2% due to double counting. We also found that 2% of individuals included in our analyses had received multiple HIV tests. These phenomena were driven by undocumented transfers. Further analysis of engagement patterns found a low level of stable engagement in care (<33%). Engagement fell into three distinct clusters: (i) characterised by high rates of late ART initiation, unstable engagement in care, and high mortality (53.9%), (ii) characterised by early ART initiation followed by prolonged periods of LTFU (13.7%), and (iii) characterised by early ART initiation followed by stable engagement in care (32.4%). Compared to cluster (i) older individuals were less likely to be in cluster (ii) and more likely to be in cluster (iii). Those who initiated ART prior to 2016 were more likely to be in cluster (ii) and (iii) compared to cluster (i). Those who initiated ART for PMTCT (RRR: 1.88 (95% CI: 1.45, 2.44)) or TB coinfection (RRR: 2.11 (95% CI: 1.27, 3.50)) were more likely to be in cluster (ii) when compared to those who initiated ART due to CD4 eligibility criteria. Males (RRR: 0.63 (95% CI: 0.51, 0.77)) were less likely to be in cluster (iii)

compared to cluster (i) as were those who initiated ART for PMTCT (RRR: 0.77 (95% CI: 0.62, 0.97)) or under test and treat guidelines when compared to those who initiated ART due to CD4 eligibility. Only a minority of patients are consistently engaged in care while the majority cycle between engagement and disengagement. Individual level data could be useful in monitoring programmes and accurately reporting patient figures if it is of high quality, has minimal missingness and is properly linked in order to account for biases that accrue from using this kind of data.

Introduction

With antiretroviral therapy (ART) programmes entering their third decade in sub-Saharan Africa (SSA), there is now a concerted effort to attain sustainable HIV epidemic control by 2030 [1–3]. Since their inception in the early 2000s, SSA ART programmes have evolved rapidly and have done a commendable job in expanding access to treatment for people living with HIV (PLHIV) [4, 5]. Expansion has been characterised by ART availability in a greater number of facilities, decentralised service delivery to bring ART closer to patients, and patient-centred care with multiple models of treatment delivery, all leading to improved convenience for patients [6]. In South Africa, the benefits of treatment are reflected in the finding that the life expectancy of PLHIV who are virally suppressed is only slightly lower than among people who are HIV-uninfected. However, PLHIV who are not virally suppressed have substantially lower life expectancy that uninfected individuals [7].

The expansion in treatment programmes has not been matched by an appropriate growth in data infrastructure, with many programmes in SSA still reliant on paper registers. Where electronic medical records (EMR) have been implemented, they are often standalone, rather than being networked across health facilities, resulting in programmes often only being able to monitor facility-based retention rather than whole programme retention [8, 9]. This limitation in monitoring individual care pathways is of concern given the importance of measuring and promoting optimal adherence to ART to achieve viral suppression, and the need to better understand patterns of engagement with HIV services, as well as transfers between these services.

In recognition that as the number of PLHIV in care increases so too does the number of care pathways, researchers have proposed a person-centred cascade that takes what some have called the "side door" into consideration [10]. This new cascade is designed to better reflect individual pathways of care including, for example, cases of loss to follow-up where individuals have reengaged at a different facility. An ability to monitor pathways of care such as this would ultimately show that ART programmes are doing better than has previously been reported [10, 11]. However, the current data infrastructure does not have the capability to do this.

In 2022, the World Health Organisation (WHO) published guidelines on strategic information for HIV advocating for a shift away from aggregated service level data to a people-centred approach supported by disaggregated longitudinal routine data [12, 13]. These guidelines included advice on patient monitoring systems, recommending the use of unique identifiers and promoting the routine collection of patient-level data for reporting on programme, national and global indicators.

In SSA advances are being made in moving towards a patient-centred approach. For example, in South Africa, efforts are underway to consolidate person-centred clinical data across government services [14]. Record linkage using demographic surveillance data has also shown

promise in improving reporting of patient outcomes in the national treatment database [15]. The ongoing transfer of paper-based clinical registers to electronic reporting, however, introduces new biases and errors. While the general direction of the potential biases has been recognised [15], there have been few efforts to ascertain the magnitude of these biases and how they may affect global targets. It is important to fill this knowledge gap as inaccurate estimates not only lead to inaccurate programme indicators but may lead to insufficient resource allocation [16], and bias projections for packages such as SPECTRUM due to erroneous modelling parameters [17].

In this study, we use public health facility ART data linked to population-based data in rural South Africa to evaluate this routine data source (health facility data) for the potential biases that might accrue when utilising it to measure programmatic indicators (e.g., numbers tested, new HIV diagnoses, numbers engaged in care) and progress towards national and global targets.

Methods

Setting and study population

Data for this study are taken from the Agincourt Health and Demographic Surveillance System (HDSS) run by the MRC/Wits Rural Public Health and Health Transitions Research Unit (Agincourt). Agincourt comprises 120,000 residents in 31 villages and is located in Mpumalanga province, South Africa where HIV prevalence was 22.8% among adults 15 to 49 years in 2017 [18–20]. Since 1992, the MRC/Wits Agincourt Research Unit has conducted an annual demographic surveillance survey where vital events (births, deaths and migration data) are collected from residents, based on a comprehensive household registration system [19, 21–23]. Since 2014, HIV patient visits to ART clinics in the area have been logged by fieldworkers and linked to the HDSS using Point-of-Contact Interactive Record Linkage (PIRL) which has been described in detail elsewhere [24, 25]. The HDSS also collects verbal autopsy data to ascertain probable causes of death [26, 27].

The study population included all ART patients aged 18 years or older, enrolled at an ART clinic (8 clinics in total) after record linkage was established in April 2014 and who had been matched to a resident record within the HDSS database. Test and treat was adopted in 2016 [28]. This cohort was followed until July 11, 2018, when data extraction occurred. De-identified clinical and demographic data used in these analyses were accessed on August 1, 2019.

Data preparation and statistical analysis

As facilities usually organise refill schedules around a 28-day schedule (2 days of extra pills), each refill was equated to one month (1 month equals 28 days). As we had no adherence data, we assumed that individuals were taking treatment as prescribed. For each eligible individual, we allocated a treatment pathway of up to 56 months (maximum follow-up in the data) from the first date that an ART facility record was opened for them (either HIV diagnosis or ART initiation). Considering the data requirements of a person-centred cascade, we assigned care pathways into seven possible states (Table 1).

We identified clusters with similar care engagement patterns using Optimal Matching distances [29, 30]. Using a binary variable coded as 1 if an individual was a member of a given cluster and 0 otherwise, factors associated with membership to each engagement-cluster were determined using multinomial logistic regression. We calculated counts and proportions for socio-demographic and baseline clinical characteristics and report confidence intervals for proportions of patients with multiple HIV tests. For continuous variables, we report the median and interquartile range. Missingness in exposure variables ranged from 4–7%. In

Table 1. Description of possible states.

State	Definition
HIV+ no ART	Either diagnosed HIV-positive but ART not initiated at one of the facilities under surveillance, or ART initiated but with no follow-up visits following initiation.
On ART	At least one health facility visit following the ART initiation visit. Individuals remained on ART until the next scheduled visit.
Late	Up to 3 months late for a scheduled ART refill visit.
Lost to follow-up (LTFU)	More than 3 months late for a scheduled ART refill visit.
Deceased	Registered as deceased from data extracted from HDSS records.
Transferred	Treatment collection from a health facility within the HDSS other than the initiating facility.
Re-engaged	Resumed treatment at initiating facility following LTFU (i.e., returned following a period of >3 months after a scheduled ART refill visit).

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logistic regression models, we only conducted complete case analyses. Sequence analyses were conducted using the *TraMineR* R-package [31]. All other analyses were conducted using Stata 16 [32]. To explore potential biases and errors we considered individuals who have tested more than once, those who have been linked to more than one clinic record, and those LTFU.

Ethical approval

Ethical approval was obtained from the London School of Hygiene and Tropical Medicine Research Ethics Committee, the University of Witwatersrand Human Research Ethics Committee, and the Mpumalanga Department of Health Public Health Research Committee. All methods were carried out in accordance with relevant guidelines and regulations and written informed consent was obtained from all participants and from a parent and/or legal guardian of minors (subjects under 18) in the study.

Results

Population characteristics

Over the study period, 8079 patient records were entered into the PIRL database. Of these, 5084 (62.9%) were linked to an HDSS resident and met the inclusion criteria. These 5084 patient records were linked to 4947 unique HDSS residents (134 unique individuals linked to multiple patient records). (S1 Fig) The 4947 individuals had a median age of 34 years (IQR: 27, 43) at ART initiation, and median baseline CD4 of 272 cells/μL (IQR: 140, 435). The 134 individuals with multiple linkages had a median age of 28 years (IQR: 24, 35), a median baseline CD4 of 268 cells/μL (IQR: 149, 425), 114 (85.1%) were female, 54 (40.3%) initiated ART in 2015, 33 (24.6%) in 2016, and 31 (23.1%) in 2014; and 63 (47.0%) initiated ART due to CD4 criteria and 38 (28.4%) for prevention of mother-to-child transmission (PMTCT). Of the 134 (2.7%) with multiple patient records, 127 (2.6%) had transferred once, 3 (0.1%) had transferred twice, and 4 (0.1%) had re-engaged at the same facility but had been given a new file number. Of 133 total transfers, 106 (79.7%) were undocumented according to PIRL records.

Based on patient records, 226 (4.6%) patients were recorded as transfers-in from facilities outside the HDSS and 47 (0.9%) were recorded as transfers-in from facilities within the HDSS at their first facility. For the 47 transfers-in from HDSS facilities, we could not identify any earlier records within PIRL, suggesting missed links or that they never had a treatment file opened at these facilities (e.g., newly diagnosed who request to start treatment at another facility).

Testing and ART initiation statistics

Of 4947 individuals, 93 [1.9% (95% CI: 1.5–2.3)] had multiple HIV tests conducted. The overwhelming majority of patients with multiple HIV tests were among those who had transferred once (i.e., linked to two records) [90/127, 70.9% (95% CI: 62.1–78.6)] and the remainder were for those who had transferred twice (i.e., linked to three records) [3/3, 100% (95% CI: 29.2–100)]. Multiple HIV tests were more common for undocumented transfers (89/106 [84.0% (95% CI: 75.6–90.4)] compared to documented transfers 4/27 [14.8% (95% CI: 4.2–33.7)]). Median time between tests was 438.5 days (IQR: 222–805).

Almost half of the 4947 individuals (46.0%; 2277) did not initiate ART immediately (i.e., within the first month) following their diagnosis. This proportion, however, declined over the study period from 48.2% (432/897)) in 2014 to 32.7% (122/373) in 2018.

Engagement sequences and clusters

The 4947 individuals contributed 147,651 person-months of engagement data to the sequence analysis corresponding to 85509 (57.9%) months in care, 47306 (32.0%) months out of care (LTFU), 11226 (7.6%) months late for a scheduled appointment, and 3610 (2.4%) for those deceased. Fig 1 illustrates the proportion of patients in each state over their engagement period (in months) by reason for ART initiation. Patients who initiated treatment for prevention of mother-to-child transmission (PMTCT) or under test and treat (strategy where persons diagnosed receive early treatment regardless of disease stage) contributed very little time [PMTCT: 161/25,949 months (0.62%); test and treat: 80/9,453 months (0.85%)] to the deceased state even later in their treatment course. Patients who initiated ART due to WHO stage (i.e., late-stage disease) had the largest contribution [466/6,937 months (6.72%)] to the deceased state.
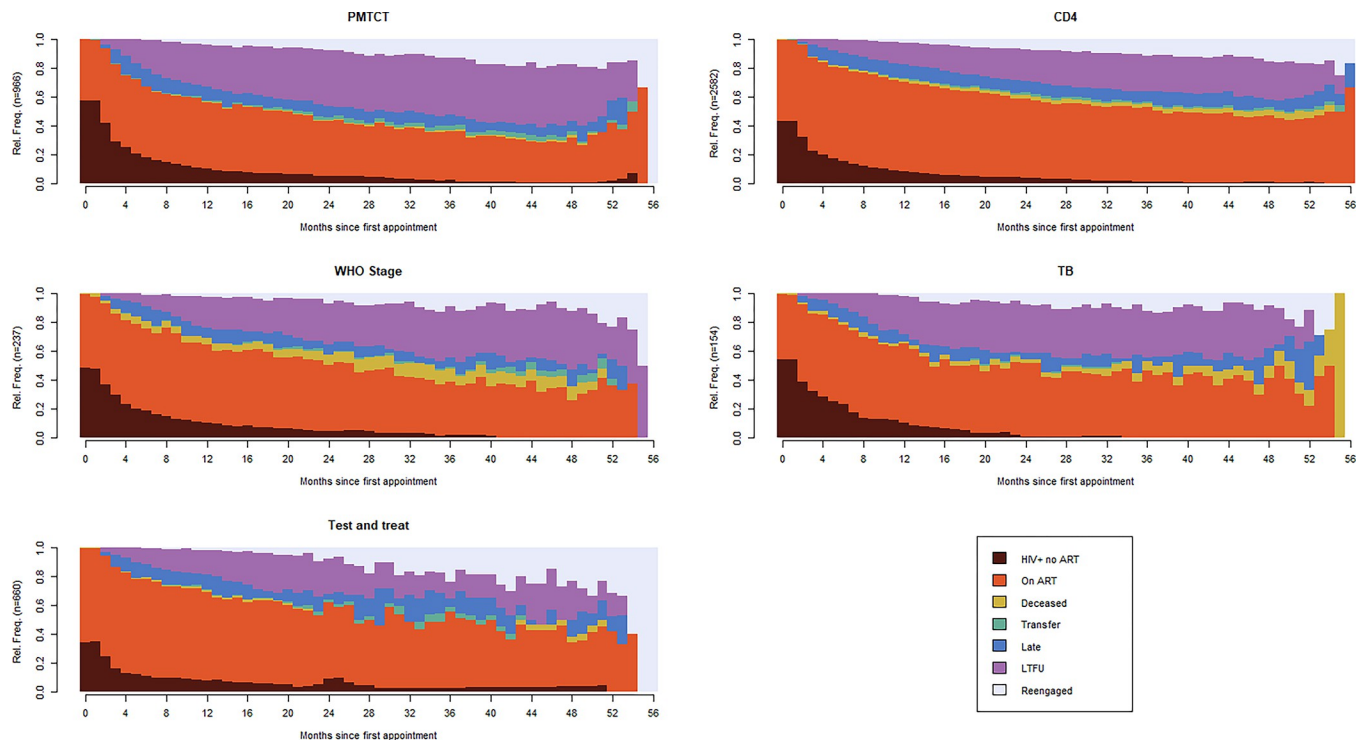

Fig 1. Chronograms of engagement in care by reason for ART initiation.

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Some patients who initiated ART for PMTCT or under test and treat also had prolonged periods where large proportions did not initiate ART (Fig 1).

An analysis of average time spent in each state shows that males were most likely to have died (males: on average 5% of sequence time versus females: 2%) as were individuals in the 55 + age group (18–24: 1%, 25–34: 2%, 35–44: 3%, 45–54: 3%, 55+: 6%), those who initiated ART with Tuberculosis (TB), or other opportunistic infections (OIs) as measured by WHO stage at ART initiation (PMTCT: 1%, CD4: 2%, WHO stage: 6%, TB: 3%, test and treat: 1%), those who initiated ART in 2014 (2014: 4%, 2015: 2%, 2016: 2%, 2017: 1%, 2018: 1%), or with lower baseline CD4 (<100: 6%, 100–199: 3%, 200–349: 2%, 350–499: 1%, 500+: 1%). Younger patients were most likely to be LTFU (18–24: 30%, 25–34: 23%, 35–44: 17%, 45–54: 15%, 55+: 10%), as were patients who initiated ART for PMTCT (PMTCT: 29%, CD4: 17%, WHO stage: 21%, TB: 24%, test and treat: 13%), and patients who initiated ART in earlier years (2014: 20%, 2015: 22%, 2016: 20%, 2017: 12%, 2018: 2%). There were no discernible differences in person-months spent in the late sequence state. (S2–S6 Figs)

We identified three distinct patterns of engagement through cluster analysis of the 4947 sequences: (i) characterised by high rates of late ART initiation, unstable engagement in care, and high mortality (n = 2666, 53.9%), (ii) characterised by early ART initiation followed by prolonged periods of LTFU (n = 679, 13.7%), and (iii) characterised by early ART initiation followed by stable engagement in care (n = 1602, 32.4%). (Table 2) Fig 2 shows individual patient trajectories further illustrating that even within the three broader engagement patterns, patient experiences of care are not monolithic. For example, for patients in the stable engagement category, there are intermittent periods of LTFU and for patients in the prolonged LTFU category, there are periods of re-engagement in care.

In the multinomial logistic regression model, older individuals (35 and older) were less likely to be in cluster 2 (early ART/prolonged LTFU) compared to cluster 1 (late ART/unstable) [e.g., RRR 55+ years versus 25–34 years: 0.44 (95% CI: 0.28, 0.72)]. Individuals who initiated ART for PMTCT when compared to those who initiated ART due to CD4 eligibility criteria were more likely to be in cluster 2 than cluster 1 [RRR: 1.88 (95% CI: 1.45, 2.44)] as were those who initiated ART due to TB coinfection. Individuals who initiated ART prior to 2016 were more likely to be in cluster 2 compared to cluster 1 [2014 vs 2016 RRR: 1.97 (95% CI: 1.49, 2.61)]. Older individuals were more likely to be in cluster 3 (early ART/stable) compared to cluster 1 as were individuals who initiated ART prior to 2016. Males were less likely to be in cluster 3 compared to cluster 1 as were individuals who initiated ART for PMTCT and those who initiated ART under test and treat (Table 3).

Discussion

In recognition that people often receive services, including HIV diagnoses, multiple times in the same or different locations, and that care pathways following a diagnosis can often be complex, WHO in 2022 recommended the collection and use of routinely linked individual data on HIV diagnoses and care to monitor people's movement over time between facilities and locations [33]. To better reflect the actual experiences of people living with diagnosed HIV, WHO also recommended the use of a new person-centred cascade. Ehrenkranz et al present a cascade that can track individuals as they cycle between disengagement and reengagement in care [11]. To achieve this individual-level data linked over time and space are required. Using a novel approach linking population-based demographic surveillance data and clinic records, we followed individuals across their treatment journey in order to assess the added value of disaggregated longitudinal patient-level data and identify and describe potential biases and errors.

Table 2. Demographic and clinical characteristics of the study cohort and engagement-cluster members.

	Demographics		Engagement clusters					
	All	%	Late ART/ Unstable	%	Early ART/Prolonged LTFU	%	Early ART/Stable	%
	4947		2666		679		1602	
Age group								
18–24	766	15.5	431	16.2	162	23.9	173	10.8
25–34	1831	37	987	37.0	296	43.6	548	34.2
35–44	1264	25.5	687	25.8	133	19.6	444	27.7
45–54	637	12.9	341	12.8	58	8.5	238	14.9
55+	449	9.1	220	8.2	30	4.4	199	12.4
Baseline CD4								
<100	868	17.6	469	17.6	116	17.1	283	17.7
100–199	869	17.6	467	17.5	104	15.3	298	18.6
200–349	1244	25.1	656	24.6	184	27.1	404	25.2
350–499	882	17.8	427	16	128	18.9	327	20.4
500+	870	17.6	491	18.4	123	18.1	256	16.0
Missing	214	4.3	156	5.9	24	3.5	34	2.1
Sex								
Female	3740	75.6	1943	72.9	517	76.1	1280	79.9
Male	1207	24.4	723	27.1	162	23.9	322	20.1
Baseline WHO stage								
I	3572	72.2	1978	74.2	464	68.3	1130	70.5
II	574	11.6	264	9.9	71	10.5	239	14.9
III	385	7.8	195	7.3	67	9.9	123	7.7
IV	56	1.1	29	1.1	7	1.0	20	1.3
Missing	360	7.3	200	7.5	70	10.3	90	5.6
ART reason								
PMTCT	966	19.5	494	18.5	213	31.4	259	16.2
CD4	2582	52.2	1200	45	297	43.7	1085	67.7
WHO stage	237	4.8	117	4.4	38	5.6	82	5.1
TB	154	3.1	76	2.9	32	4.7	46	2.9
Test and Treat	660	13.4	576	21.6	26	3.8	58	3.6
Missing	348	7.0	203	7.6	73	10.8	72	4.5
ART year								
2014	897	18.1	244	9.2	162	23.9	491	30.7
2015	1246	25.2	331	12.4	277	40.8	638	39.8
2016	1395	28.2	697	26.1	225	33.1	473	29.5
2017	1036	20.9	1021	38.3	15	2.2	0	0.0
2018	373	7.5	373	14.0	0	0.0	0	0.0

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In our cohort of South African patients, we found 2% inflation of patient numbers due to double counting as a result of undocumented transfer. Of the 2.7% of people transferring between facilities during the study period, 80% of these were undocumented. We also report 2% of individuals having multiple positive tests, with individuals with undocumented transfers more likely to have multiple HIV diagnoses. HIV testing data in this setting has not been digitised and we relied heavily on tests that had been recorded in patient records. As such, this is likely an underestimation of retesting in this setting. This is especially pertinent as one study where testing data was digitised and linked to individuals in the Western Cape of South Africa found that 51% of positive diagnoses were from individuals previously diagnosed [34]. As
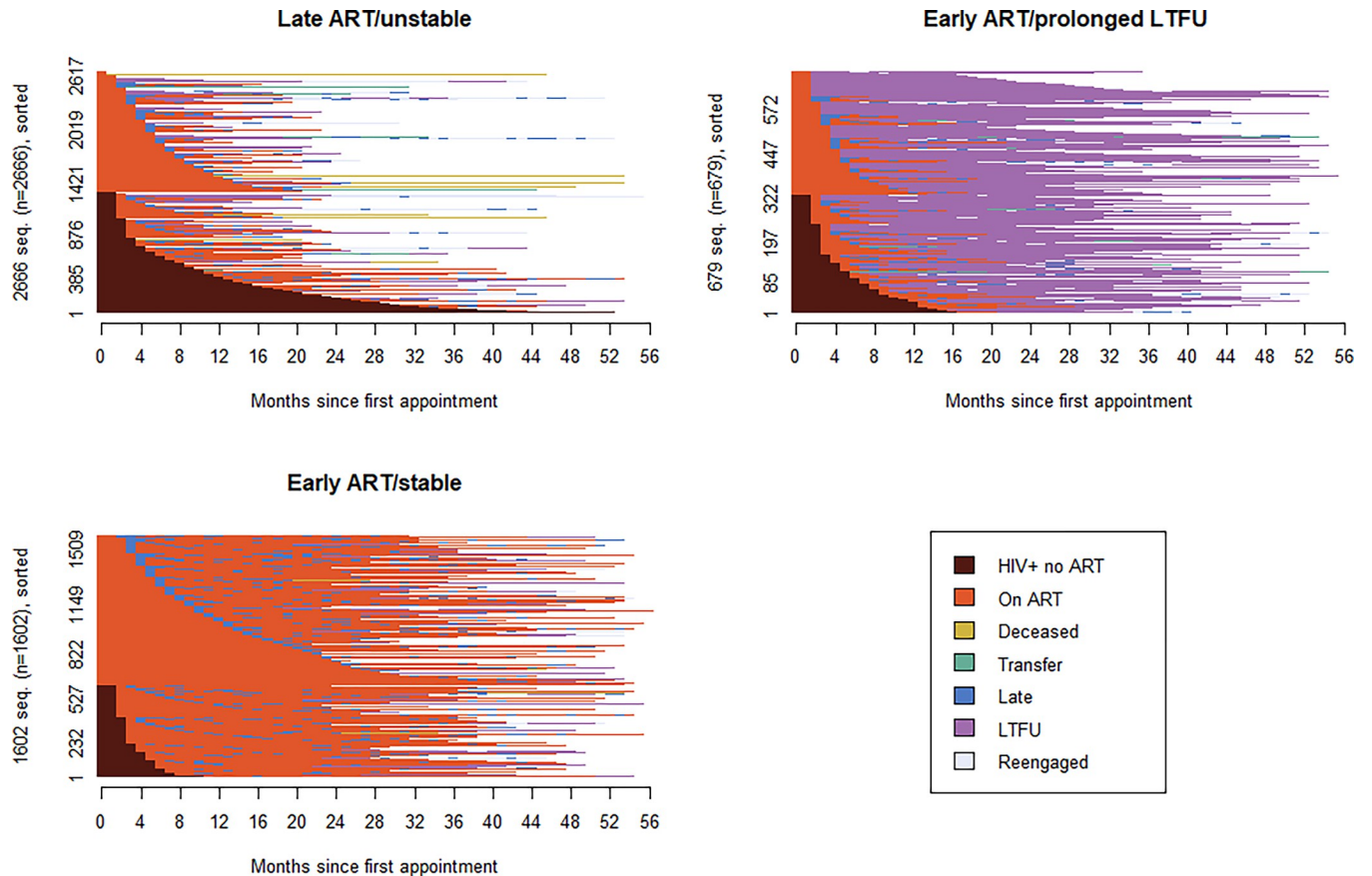
Fig 2. Index plots of the three engagement clusters showing individual patient trajectories.

ART is increasingly offered to healthier patients, undocumented transfers are becoming a more common occurrence in treatment programmes likely linked to labour migration [35], illustrating the need for robust cascades that takes into consideration, movements between clinics and regions and non-linear care pathways.

Using analysis of patient engagement sequences, we identified low levels (<33%) of stable engagement in care, categorising three distinct patterns of engagement. These were (1) high rates of late ART initiation, unstable engagement in care, and high mortality, (2) early ART initiation followed by prolonged periods of LTFU, and (3) early ART initiation followed by stable engagement in care. Historically, HIV programmes have used aggregate data which lacks the nuance of individual level data. The cluster analysis demonstrates the multiple ways that individuals engage with care showing that this does not follow a linear path. This is something that would not be possible with aggregate data. The clusters could also potentially assist programmes in estimating how many individuals within their programmes might have similar HIV care engagement profiles. We found that early ART initiation is still lagging, similar to other studies [36] and could reflect several reasons including individual readiness. Additionally, TB coinfection is common in this setting and delayed ART is recommended in such cases [37]. Individuals who were pregnant or who had OIs were more likely to initiate ART early consistent with treatment recommendations. However, these groups were likely to experience prolonged LTFU similar to findings from other studies [38]. Our findings suggest that two thirds of the treatment cohort have cyclical engagement patterns. We saw that LTFU is not a

Table 3. Factors associated with engagement-cluster membership.

	Early ART/Prolonged LTFU (n = 679) vs Late ART/Unstable		Early ART/Stable (n = 1,602) vs Late ART/Unstable	
	RRR (95% CI)	p-value	RRR (95% CI)	p-value
Age group				
18–24	1.42 (1.08, 1.87)	0.012	0.84 (0.65, 1.08)	0.18
25–34	Reference	__	Reference	__
35–44	0.75 (0.57, 0.99)	0.04	1.29 (1.05, 1.60)	0.016
45–54	0.63 (0.43, 0.92)	0.017	1.29 (0.99, 1.68)	0.057
55+	0.44 (0.28, 0.72)	0.001	1.38 (1.03, 1.86)	0.03
Sex				
Female	Reference	__	Reference	__
Male	1.23 (0.95, 1.59)	0.123	0.63 (0.51, 0.77)	<0.001
ART reason				
PMTCT	1.88 (1.45, 2.44)	<0.001	0.77 (0.62, 0.97)	0.024
CD4	Reference	__	Reference	__
WHO stage	1.24 (0.81, 1.89)	0.322	0.75 (0.53, 1.05)	0.095
TB	2.11 (1.27, 3.50)	0.004	0.87 (0.54, 1.38)	0.55
Test and Treat	0.73 (0.47, 1.16)	0.186	0.51 (0.36, 0.72)	<0.001
ART year				
2014	1.97 (1.49, 2.61)	<0.001	2.95 (2.39, 3.63)	<0.001
2015	2.61 (2.06, 3.30)	<0.001	2.79 (2.31, 3.36)	<0.001
2016	Reference	__	Reference	__
2017	0.04 (0.02, 0.07)	<0.001	Omitted	__
2018	Omitted	__	Omitted	__

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terminal state, and individuals move between periods of engagement and disengagement, something that only patient-centred data can capture. We also found that engagement sequences are as unique as the individual, as such, programmes will need to evolve to offer tailored care to fit patients' individual needs.

Although we were able to link individuals across facilities, we report there being a large proportion of individuals who remain LTFU from this network. This likely speaks to two points. Firstly, the South African population is highly mobile [39], a consequence of historic inequality where economic opportunities were only available in select parts of the country. We have shown previously that individuals from this region transferred their care as far as Cape Town, in the Western Cape [35]. Secondly, ascertainment of deaths is still difficult even with demographic surveillance data. Given that membership in the prolonged LTFU category was associated with TB coinfection as the ART initiation reason (a factor associated with high mortality in other cohorts [40, 41]), it is likely that some of these individuals might have died. Our findings illustrate that there is still a need for routine reporting mechanisms that accurately document mortality a gap also recognised by WHO, which recommends cross referencing other data sources where deaths are recorded to improve accuracy of this information [33]. We have reported previously that LTFU misclassifies a large proportion of patient outcomes [15]. As such, ascertainment of outcomes following LTFU will be important in order to more accurately estimate programmatic metrics. We should note that we were unable to link 37% of patients who claimed residency in the HDSS which might have resolved some of the LTFU.

Our study has some limitations. Linkage errors, especially due to missed matches, may bias our findings. Our linkage covered a small group of public health facilities offering ART in a rural community and did not extend to several private facilities that offer services to the community or to public facilities in surrounding areas where some members of the community

could be receiving care. More extensive geographical linkage, and the inclusion of public and private facilities could reveal higher proportions of retesting and silent transfers, and as such our estimates should be interpreted as lower limits for adjustment factors for estimates from routine programme data. Due to the routine nature of data, some variables had a high level of missing values. As we did not have adherence data, the assumption that individuals were in care between treatment refills might misclassify some individuals. Optimal matching has some limitations similar to other statistical models used for cluster analysis (e.g., a statistical technique cannot speak to individual lived experiences).

In our setting, we not only highlight the feasibility and utility of linking individual level data across time and space but also report fairly modest frequency of retesting and double counting of patients on ART, highlighting the importance of silent transfers in increasing the occurrence of these phenomena. Our findings suggest that HIV programmes in SSA may need to pay particular attention to younger individuals, pregnant women, individuals with OIs, those with higher CD4 counts at diagnosis, and males as these groups are least likely to remain stable on ART and could benefit from extra interventions.

To better understand and improve programme performance, we need to strengthen data quality, sustainability and use through precise analysis and the characterisation and correction of potential biases. With the aspiration of populating provincial and national level person-centred HIV care cascades, we need to expand the reach of data linkage across facilities to more accurately describe individual level pathways of care, thereby minimising potential biases and errors.

Supporting information

S1 Checklist. Inclusivity in global research.
(DOCX)

S1 Fig. Flow chart of individuals included in the analysis.
(DOCX)

S2 Fig. Proportion of sequence time spent in each state on average by age.
(DOCX)

S3 Fig. Proportion of sequence time spent in each state on average by ART initiation reason.
(DOCX)

S4 Fig. Proportion of sequence time spent in each state on average by baseline CD4.
(DOCX)

S5 Fig. Proportion of sequence time spent in each state on average by sex.
(DOCX)

S6 Fig. Proportion of sequence time spent in each state on average by year of ART initiation.
(DOCX)

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Conceptualization: David Etoori, Alison Wringe, Georges Reniers.

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References

1. UNAIDS. 90-90-90: An Ambitious Treatment Target to Help End the AIDS Epidemic. [Internet]. 2014 [cited 2016 Sep 16]. Available from: http://www.unaids.org/sites/default/files/media_asset/90-90-90_en_0.pdf

2. Sidibé M, Loures L, Samb B. The UNAIDS 90–90–90 target: a clear choice for ending AIDS and for sustainable health and development. J Int AIDS Soc. 2016; 19(1):21133. https://doi.org/10.7448/IAS.19.1.21133 PMID: 27424601

3. UNAIDS. Miles to go: closing gaps, breaking barriers, righting injustices [Internet]. 2018 [cited 2022 Jun 18]. Available from: https://www.unaids.org/sites/default/files/media_asset/miles-to-go_en.pdf

4. Weller I. Delivery of Antiretroviral Therapy in Sub-Saharan Africa. Clin Infect Dis. 2006 Sep 15; 43 (6):777–8. https://doi.org/10.1086/507113 PMID: 16912955

5. Ford N, Calmy A, Mills EJ. The first decade of antiretroviral therapy in Africa. Glob Health. 2011 Sep 29; 7(1):33. https://doi.org/10.1186/1744-8603-7-33 PMID: 21958478

6. Rabkin M, El-Sadr WM, De Cock KM, Bellagio HIV/Health Systems Working Group1. The impact of HIV scale-up on health systems: A priority research agenda. J Acquir Immune Defic Syndr 1999. 2009 Nov; 52 Suppl 1:S6–11. https://doi.org/10.1097/QAI.0b013e3181bbcd69 PMID: 19858943

7. Payne CF, Houle B, Chinogurei C, Herl CR, Kabudula CW, Kobayashi LC, et al. Differences in healthy longevity by HIV status and viral load among older South African adults: an observational cohort modelling study. Lancet HIV. 2022 Oct 1; 9(10):e709–16. https://doi.org/10.1016/S2352-3018(22)00198-9 PMID: 36179754

8. Gourlay A, Wringe A, Todd J, Michael D, Reniers G, Urassa M, et al. Challenges with routine data sources for PMTCT programme monitoring in East Africa: insights from Tanzania. Glob Health Action. 2015; 8:29987. https://doi.org/10.3402/gha.v8.29987 PMID: 26715204

9. Etoori D, Wringe A, Renju J, Kabudula CW, Gomez-Olive FX, Reniers G. Challenges with tracing patients on antiretroviral therapy who are late for clinic appointments in rural South Africa and recommendations for future practice. Glob Health Action [Internet]. 2020 Apr 28 [cited 2020 Jul 31]; 13(1). Available from: https://www.ncbi.nlm.nih.gov/pmc/articles/PMC7241554/ https://doi.org/10.1080/16549716.2020.1755115 PMID: 32340584

10. Hallett TB, Eaton JW. A side door into care cascade for HIV-infected patients? J Acquir Immune Defic Syndr 1999. 2013 Jul; 63 Suppl 2:S228–232. https://doi.org/10.1097/QAI.0b013e318298721b PMID: 23764640

11. Ehrenkranz P, Rosen S, Boulle A, Eaton JW, Ford N, Fox MP, et al. The revolving door of HIV care: Revising the service delivery cascade to achieve the UNAIDS 95-95-95 goals. PLoS Med. 2021 May 24; 18(5):e1003651. https://doi.org/10.1371/journal.pmed.1003651 PMID: 34029346

12. WHO. Consolidated guidelines on the use of antiretroviral drugs for treating and preventing HIV infection Recommendations for a public health approach [Internet]. 2016. Available from: http://www.who.int/hiv/pub/arv/arv-2016/en/

13. World Health Organisation. Consolidated guidelines on person-centred HIV patient monitoring and case surveillance [Internet]. 2017. Available from: https://www.who.int/hiv/pub/guidelines/person-centred-hiv-monitoring-guidelines/en/

14. Boulle A, Vega I de, Moodley M, Shand L, Loff A, Fredericks 004E, et al. Data Centre Profile: The Provincial Health Data Centre of the Western Cape Province, South Africa. Int J Popul Data Sci [Internet]. 2019 Nov 20 [cited 2023 Jun 30]; 4(2). Available from: https://ijpds.org/article/view/1143 https://doi.org/10.23889/ijpds.v4i2.1143 PMID: 32935043

15. Etoori D, Wringe A, Kabudula CW, Renju J, Rice B, Gomez-Olive FX, et al. Misreporting of patient outcomes in the South African national HIV treatment database: consequences for programme planning, monitoring, and evaluation. Front Public Health [Internet]. 2020 [cited 2020 Mar 24]; 8. Available from: https://www.frontiersin.org/articles/10.3389/fpubh.2020.00100/full#supplementary-material

16. Mate KS, Bennett B, Mphatswe W, Barker P, Rollins N. Challenges for routine health system data management in a large public programme to prevent mother-to-child HIV transmission in South Africa. PloS One. 2009; 4(5):e5483. https://doi.org/10.1371/journal.pone.0005483 PMID: 19434234

17. Ghys PD, Brown T, Grassly NC, Garnett G, Stanecki KA, Stover J, et al. The UNAIDS Estimation and Projection Package: a software package to estimate and project national HIV epidemics. Sex Transm Infect. 2004 Aug 1; 80(suppl 1):i5–9. https://doi.org/10.1136/sti.2004.010199 PMID: 15249692

18. Human Sciences Research Council (HSRC). The fifth South African national HIV prevalence, incidence, behaviour and communication survey, 2017 (SABSSM V) [Internet]. 2018 [cited 2021 Nov 13]. Available from: http://www.hsrc.ac.za/uploads/pageContent/9234/SABSSMV_Impact_Assessment_Summary_ZA_ADS_cleared_PDFA4.pdf

19. Kahn K, Collinson MA, Gómez-Olivé FX, Mokoena O, Twine R, Mee P, et al. Profile: Agincourt health and socio-demographic surveillance system. Int J Epidemiol. 2012 Aug; 41(4):988–1001. https://doi.org/10.1093/ije/dys115 PMID: 22933647

20. Kabudula CW, Houle B, Collinson MA, Kahn K, Tollman S, Clark S. Assessing Changes in Household Socioeconomic Status in Rural South Africa, 2001–2013: A Distributional Analysis Using Household Asset Indicators. Soc Indic Res. 2016 Jun 28;1–27. https://doi.org/10.1007/s11205-016-1397-z PMID: 28931968

21. Tollman SM. The Agincourt field site—evolution and current status. South Afr Med J Suid-Afr Tydskr Vir Geneeskd. 1999 Aug; 89(8):853–8. PMID: 10488361

22. Tollman SM, Herbst K, Garenne M, Gear JS, Kahn K. The Agincourt demographic and health study—site description, baseline findings and implications. South Afr Med J Suid-Afr Tydskr Vir Geneeskd. 1999 Aug; 89(8):858–64. PMID: 10488362

23. Tollman S, Kahn K. Report on the meeting 'Strengthening ties: the Agincourt field site in its African context.' Trop Med Int Health. 1997; 2(9):920–3.

24. Rentsch CT, Kabudula CW, Catlett J, et al. Point-of-contact Interactive Record Linkage (PIRL): A software tool to prospectively link demographic surveillance and health facility data. Gates Open Res [Internet]. 2017; 1(8). Available from: https://gatesopenresearch.org/articles/1-8/v1 https://doi.org/10.12688/gatesopenres.12751.2 PMID: 29528050

25. Kabudula CW, Clark BD, Gómez-Olivé FX, Tollman S, Menken J, Reniers G. The promise of record linkage for assessing the uptake of health services in resource constrained settings: a pilot study from South Africa. BMC Med Res Methodol. 2014 May 24; 14:71. https://doi.org/10.1186/1471-2288-14-71 PMID: 24884457

26. Byass P, Hussain-Alkhateeb L, D'Ambruoso L, Clark S, Davies J, Fottrell E, et al. An integrated approach to processing WHO-2016 verbal autopsy data: the InterVA-5 model. BMC Med. 2019 30; 17 (1):102. https://doi.org/10.1186/s12916-019-1333-6 PMID: 31146736

27. Kabudula CW, Houle B, Collinson MA, Kahn K, Gómez-Olivé FX, Tollman S, et al. Socioeconomic differences in mortality in the antiretroviral therapy era in Agincourt, rural South Africa, 2001–13: a population surveillance analysis. Lancet Glob Health. 2017; 5(9):e924–35. https://doi.org/10.1016/S2214-109X(17)30297-8 PMID: 28807190

28. South Africa: National Department of Health. Implementation of the universal test and treat strategy for HIV positive patients and differentiated care for stable patients [Internet]. 2016 [cited 2019 Jul 24]. Available from: https://sahivsoc.org/Files/22%208%2016%20Circular%20UTT%20%20%20Decongestion%20CCMT%20Directorate.pdf

29. Abbott A, Forrest J. Optimal Matching Methods for Historical Sequences. J Interdiscip Hist. 1986; 16 (3):471.

30. Abbott A, Tsay A. Sequence Analysis and Optimal Matching Methods in Sociology: Review and Prospect. Sociol Methods Res. 2000 Aug 1; 29(1):3–33.

31. Gabadinho A, Ritschard G, Müller NS, Studer M. Analyzing and Visualizing State Sequences in R with TraMineR. J Stat Softw. 2011 Apr 7; 40(1):1–37.

32. StataCorp. Stata Statistical Software. College Station, TX: StataCorp LLC; 2017.

33. WHO. Consolidated guidelines on person-centred HIV strategic information: strengthening routine data for impact [Internet]. 2022 [cited 2023 Jun 30]. Available from: https://www.who.int/publications-detail-redirect/9789240055315

34. Jacob N, Rice B, Kalk E, Heekes A, Morgan J, Hargreaves J, et al. Utility of digitising point of care HIV test results to accurately measure, and improve performance towards, the UNAIDS 90-90-90 targets. PLOS ONE. 2020 Jun 30; 15(6):e0235471. https://doi.org/10.1371/journal.pone.0235471 PMID: 32603374

35. Etoori D, Kabudula CW, Wringe A, Rice B, Renju J, Gomez-Olive FX, et al. Investigating clinic transfers among HIV patients considered lost to follow-up to improve understanding of the HIV care cascade: Findings from a cohort study in rural north-eastern South Africa. PLOS Glob Public Health. 2022 May 24; 2(5):e0000296. https://doi.org/10.1371/journal.pgph.0000296 PMID: 36962304

36. Lilian RR, Rees K, McIntyre JA, Struthers HE, Peters RPH. Same-day antiretroviral therapy initiation for HIV-infected adults in South Africa: Analysis of routine data. PLoS ONE. 2020 Jan 14; 15(1):e0227572. https://doi.org/10.1371/journal.pone.0227572 PMID: 31935240

37. Meintjes G, Moorhouse MA, Carmona S, Davies N, Dlamini S, van Vuuren C, et al. Adult antiretroviral therapy guidelines 2017. South Afr J HIV Med. 2017; 18(1):776. https://doi.org/10.4102/sajhivmed.v18i1.776 PMID: 29568644

38. Knettel BA, Cichowitz C, Ngocho JS, Knippler ET, Chumba LN, Mmbaga BT, et al. Retention in HIV Care During Pregnancy and the Postpartum Period in the Option B+ Era: A Systematic Review and Meta-Analysis of Studies in Africa. JAIDS J Acquir Immune Defic Syndr [Internet]. 2017 Dec 4 [cited 2018 Jan 7];Publish Ahead of Print. Available from: http://journals.lww.com/jaids/Abstract/publishahead/Retention_in_HIV_Care_During_Pregnancy_and_the.96776.aspx

39. Collinson MA. Striving against adversity: the dynamics of migration, health and poverty in rural South Africa. Glob Health Action. 2010 Jun 3; 3.

40. Hunt PW, Lee SA, Siedner MJ. Immunologic Biomarkers, Morbidity, and Mortality in Treated HIV Infection. J Infect Dis. 2016 Oct 1; 214(Suppl 2):S44–50. https://doi.org/10.1093/infdis/jiw275 PMID: 27625430

41. Lelisho ME, Wotale TW, Tareke SA, Alemu BD, Hassen SS, Yemane DM, et al. Survival rate and predictors of mortality among TB/HIV co-infected adult patients: retrospective cohort study. Sci Rep. 2022 Nov 1; 12(1):18360. https://doi.org/10.1038/s41598-022-23316-4 PMID: 36319734