

Transmission and case projections: 2000 to 2050

Data and analysis tool (GUI)

Gambiense Human African Trypanosomiasis (gHAT or sleeping sickness) disease transmission model - **data and results**

User Guide

Contents

[Quick guide to getting started](#)

[Definitions](#)

[Results tabs](#)

❖ [Screening](#)

❖ [Detections & New Infections](#)

○ [Prospective Model Analysis](#)

○ [Retrospective Model Analysis](#)

ABOUT

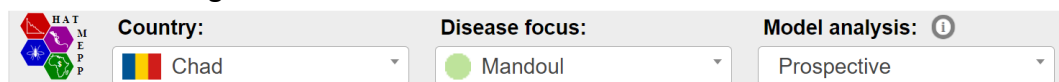
DESCRIPTION	gHAT transmission and case projections
SOURCE	Projections from Warwick gHAT model fitted to <u>WHO HAT Atlas</u> case data (2000-2018) and PNLTHA-Chad data 2019
DATE ISSUED	May 2021
LAST UPDATED	April 2022
SPATIAL COVERAGE	Mandoul, Chad
YEARS(S)	Fitting to 2000 to 2019 and projections from 2020 to 2050
STATUS	Completed
CREATOR	Rock et al, University of Warwick
USAGE	Open access
CONTACT EMAIL	<u>K.S.Rock@warwick.ac.uk</u>

Quick guide to getting started

- 1) Go to: <https://hatmepp.warwick.ac.uk/Mandoulfitandproject/v1/index.php>

(We recommend you use Google Chrome, Microsoft Edge or Firefox as your browser to get the best experience)

- 2) Choose your **disease focus** and **model analysis** from the drop-down boxes. This version of the GUI is only for Chad and Mandoul so the country and disease focus cannot be changed.



The screenshot shows a user interface with three main sections. On the left is a logo for HATMEPP (Health, Antigen, and Transmission Modelling and Epidemiology Project) consisting of four colored circles (red, purple, green, blue) with icons inside. To the right of the logo is a 'Country:' dropdown menu showing the flag of Chad and the text 'Chad'. Further right is a 'Disease focus:' dropdown menu with a green circle icon and the text 'Mandoul'. On the far right is a 'Model analysis:' dropdown menu with an information icon and the text 'Prospective'.

Note that default results are for Mandoul and the Prospective model analysis

- 3) The table immediately below will auto generate based on your entries, to show the disease focus population, proportion of people assumed to be screened, deployment of vector control and year of improved medical interventions.



Chad: Mandoul disease focus	
Information	
Chad population (2017)	14,899,994
Life expectancy in 2017 (F/M)	61.7/58.6
Mandoul population (est 2015)	41,000
Screening level (mean/max)	48%/67%
Vector control	Begun 2014, reduction 99.1% [95.8%, 99.6%]
Improved medical	Begun 2015

- 4) Several [results tabs](#) can be found under the maps and table. **Charts** under each results tab will auto-generate based on your disease focus.
- 5) You can **download charts**, by clicking on 'Save Plot' (bottom left of each screen).

Definitions

<i>Terminology</i>	<i>Definition</i>
<i>Assumed (max)</i>	Assumed number of people screened in the projections in the selected foci. This is based on the historical maximum level of active screening (see <i>Max AS below</i>)
<i>Assumed (mean)</i>	Assumed number of people screened in the projections in the selected foci under a mean level of active screening (see <i>Mean AS below</i>)
<i>Fitted</i>	Model outputs have been fitted to actual case data from the WHO HAT Atlas (2000-2018) and PNLTHA-Chad (2019)
<i>Mean AS</i>	The proportion of people screened is equal to the mean number screened during 2015–2019
<i>Max AS</i>	The proportion of people screened is equal to the historical maximum number screened during 2000–2019
<i>Observed</i>	Aggregate case data from the WHO HAT Atlas/PNLTHA-Chad
<i>PS (Passive Screening)</i>	Passive screening is in place under all strategies
<i>Improved Passive Screening (RDT)</i>	Fortification of the passive screening system by increasing the number of fixed health facilities with rapid diagnostic tests
<i>VC (Vector Control)</i>	Vector control (VC) is simulated to reflect the Tiny Target intervention which began in 2014
<i>Spec (Specificity)</i>	The proportion of people without disease that test negative
<i>Prospective (Future) Model Analysis</i>	Given the current situation the prospective model analysis examines what would be expected to happen in the future
<i>Retrospective (Past) Model Analysis</i>	Examines what would have happened had different strategies been implemented from 2014
<i>Stop 2021</i>	Active screening and vector control cessation from 2021

You can also refer to the main [Glossary](#) for a description of commonly used terms and acronyms associated with the gHAT projects.

Results tabs

[Screening Data](#)

[Active Detections](#)

[Passive Detections](#)

[New Infections](#)

Screening

Screening Data

Active Detections

Passive Detections

New Infections

The **Screening** results tab provides a chart showing you (i) the number of people actively screened by year from 2000-2019 (i.e. the “observed” level in the data) and (ii) the assumed number of people that are screened in the selected foci under a mean level of active screening compared to the historical maximum level of active screening (see [definitions](#) for mean AS and max AS) from 2019 onwards.

Display year range

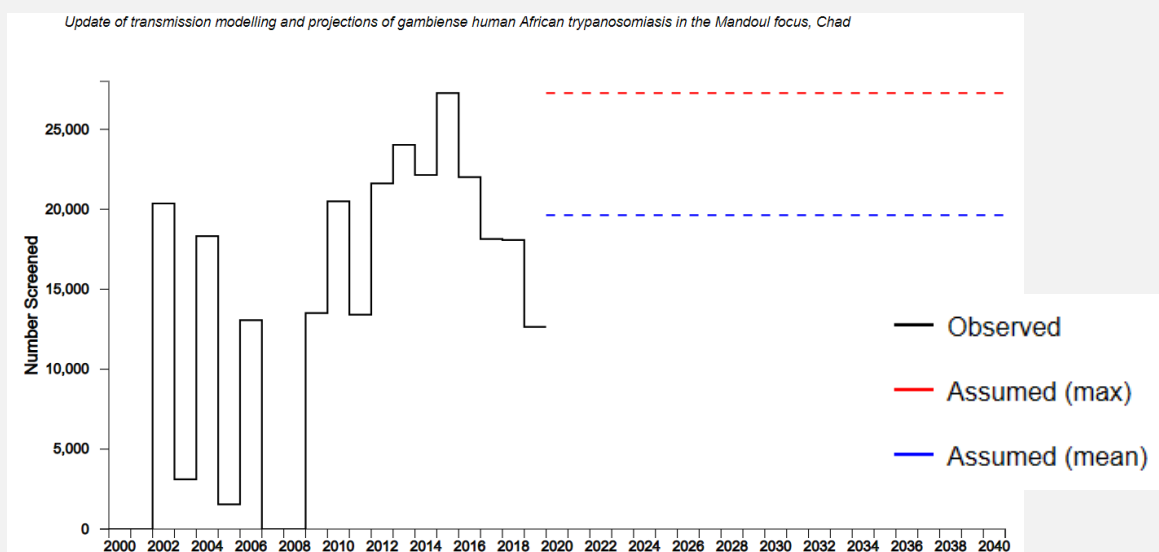
From: 2010

To: 2040

Amend the time period as required (defaulted to 2010 to 2040, but this can be scaled to any period between 2000 and 2050).

Note: Screening data remains the same irrespective of the selected model analysis

Illustration 1: Assumed number of people to be screened under a mean and maximum level of active screening



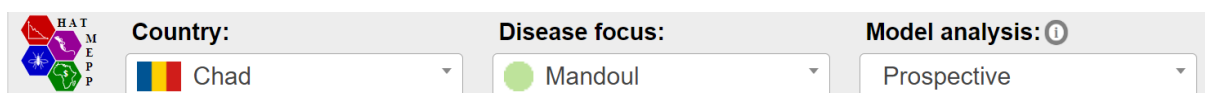
Detections & New Infections: Prospective & Retrospective Dataset Model Analysis



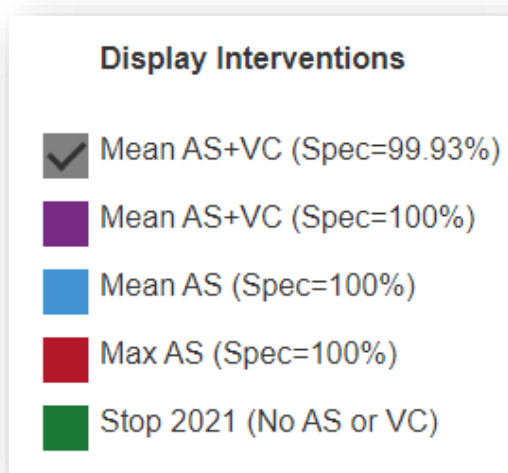
The final three results tabs provide charts to show you the predicted active and passive case reporting by year and by foci, as well as new infections that are not directly measurable, viewable for each intervention strategy (Display Interventions). The **prospective model analysis** examines the current situation in Chad (2020) and the potential impact of future interventions on detections and transmission. The **retrospective model analysis** examines what would have happened had different strategies be implemented from 2014 and how these shape future detections and transmission.

Prospective Model Analysis:

From the grey bar at the top of the page select the disease focus and prospective dataset (hover over the (i) for prospective model analysis details):



Then select the active detections, passive detections, or new infections tab below:



The list of **Display Interventions** (see opposite) appears as a tick box function to allow you to select and compare the results based on different intervention strategies. Specificity (Spec) of the diagnostic algorithm can be set at 99.93% or 100% (see [definitions](#)). The tick box will default to Mean AS + VC (Spec = 99.93%). Stop 2021 refers to the cessation of active screening and vector control activities from 2021.

Tips:

- Amend the time period as required (defaulted to 2010 to 2040, but this can be scaled to any period between 2000 and 2050).
- Hover on the results for the year you are interested in to view an information box confirming the predicted highest (97.5 percentile), lowest (2.5 percentile) and median number of cases within the range.

Example:

1. Selected foci: Mandoul from the top bar.
2. Selected dataset: Prospective model analysis from the top bar.
3. Click on the “Active Detections” tab. The range will be from 2010-2040 by default.
4. Select 2012 in the “From” dropdown list on the left-hand side and 2035 in the “To” dropdown list.
5. Show all five strategies from the article by additionally selecting Mean AS + VC (Spec = 99.93%), Mean AS + VC (Spec = 100%), Mean AS + no VC (Spec = 100%), Max AS + no VC (Spec = 100%), Stop 2021 (No AS or VC) in the right-hand side legend.
6. You can save this image by clicking “Save Plot” on the left-hand side.
7. To view predictions for Passive Detections and New Infections with these same settings, simply choose the corresponding tab. The selected strategies and data range will remain the same.

Illustration 2: Example results – predicted active detections and passive detections under five prospective intervention strategies from 2012 to 2035

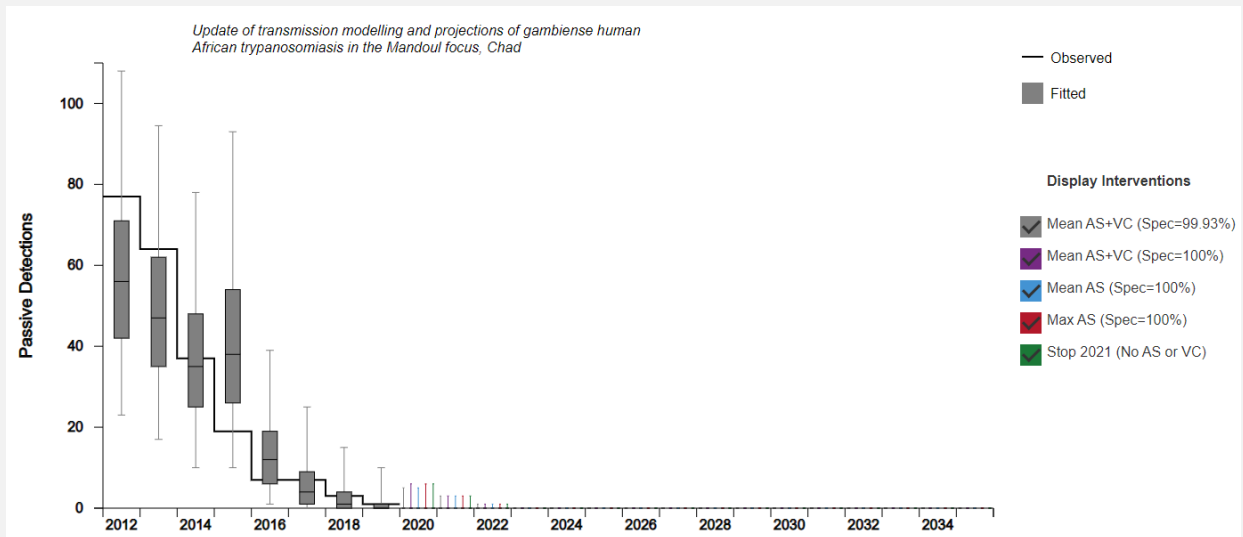
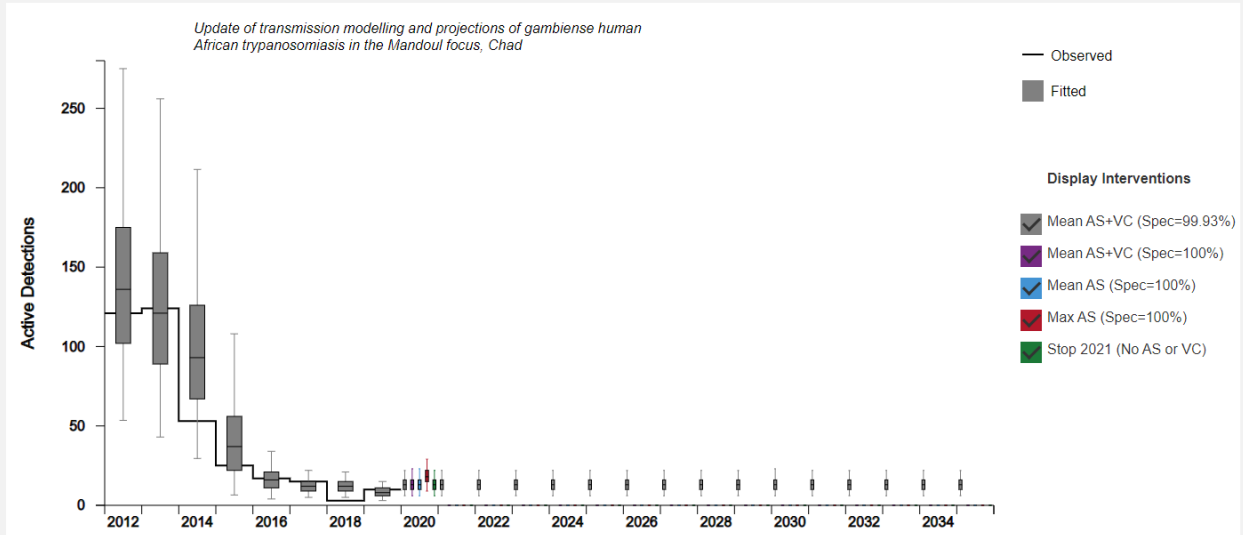
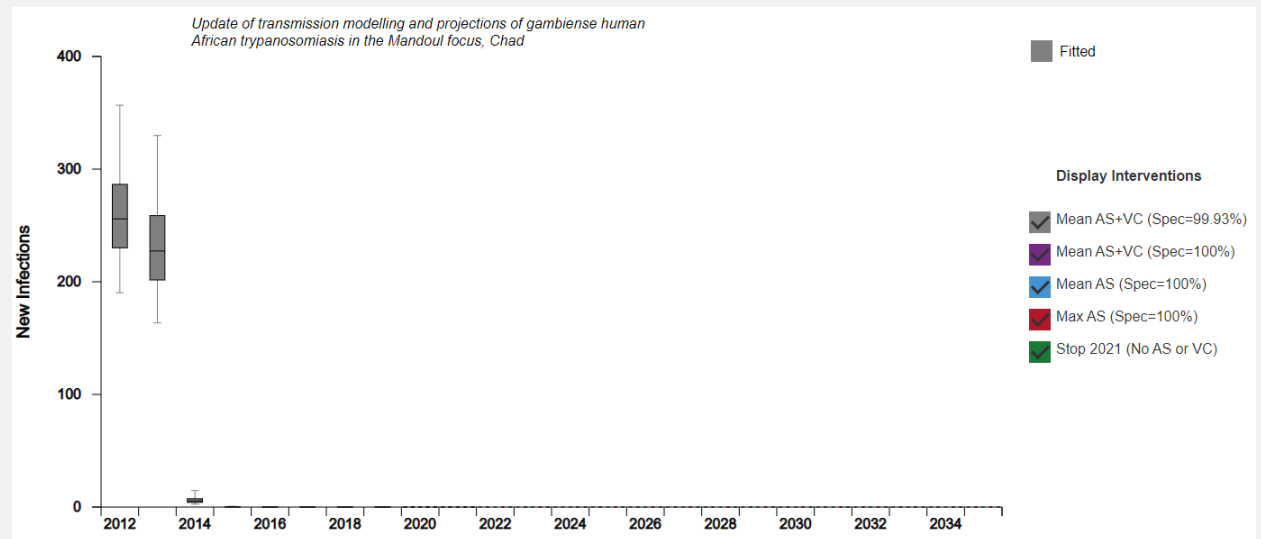
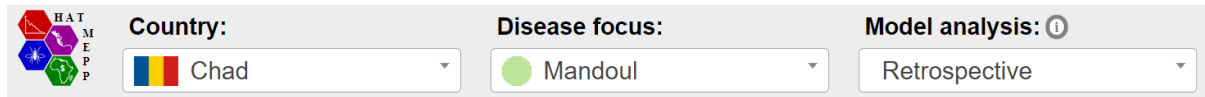


Illustration 3: Example results – predicted new infections under five prospective intervention strategies from 2012 to 2035



Retrospective Model Analysis:

From the grey bar at the top of the page select the disease focus and retrospective dataset (hover over the (i) for retrospective model analysis details):

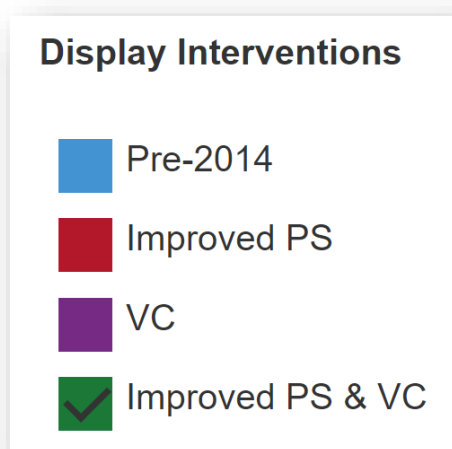


The screenshot shows a grey navigation bar with three dropdown menus. On the left is a logo with the letters 'H A T' and 'M E P P' around a central star. The first dropdown is labeled 'Country:' and shows the flag of Chad with the text 'Chad'. The second dropdown is labeled 'Disease focus:' and shows a green circle with the text 'Mandoul'. The third dropdown is labeled 'Model analysis: (i)' and shows the text 'Retrospective'.

Then select the active detections, passive detections or new infections tab below:



The screenshot shows four navigation tabs. The first tab is 'Screening Data' and is grey. The other three tabs are 'Active Detections', 'Passive Detections', and 'New Infections', all of which are blue.



The screenshot shows a white box titled 'Display Interventions'. It contains four items, each with a colored square and a text label: a blue square for 'Pre-2014', a red square for 'Improved PS', a purple square for 'VC', and a green square with a checkmark for 'Improved PS & VC'.

The list of Display Interventions (see opposite) appears as a tick box function to allow you to select and compare the results based on different intervention strategies. The tick-box will default to improved passive screening & vector control as was implemented in Chad from 2014 onwards. Pre-2014 refers to those interventions in place before improvements in passive screening and the introduction of vector control.

Tips:

- Amend the time period as required (defaulted to 2010 to 2040, but this can be scaled to any period between 2000 and 2050).
- Hover on the results for the year you are interested in to view an information box confirming the predicted highest (97.5), lowest (2.5) and median number of cases within the range.

Example:

1. Selected foci: Mandoul from the top bar.
2. Selected dataset: Retrospective dataset from the top bar.
3. Click on the “Active Detections” tab. The range will be from 2010-2040 by default.
4. Select 2012 in the “From” dropdown list on the left-hand side and 2035 in the “To” dropdown list.

5. Show all four strategies from the article by additionally selecting pre-2014, improved PS, VC and Improved PS & VC I in the right-hand side legend.
6. You can save this image by clicking “Save Plot” on the left-hand side.
7. To view predictions for Passive Detections and New Infections with these same settings, simply choose the corresponding tab. The selected strategies and data range will remain the same until you choose a new foci.

Illustration 4: Example results – predicted active detections and passive detections under four retrospective intervention strategies (2012 to 2035)

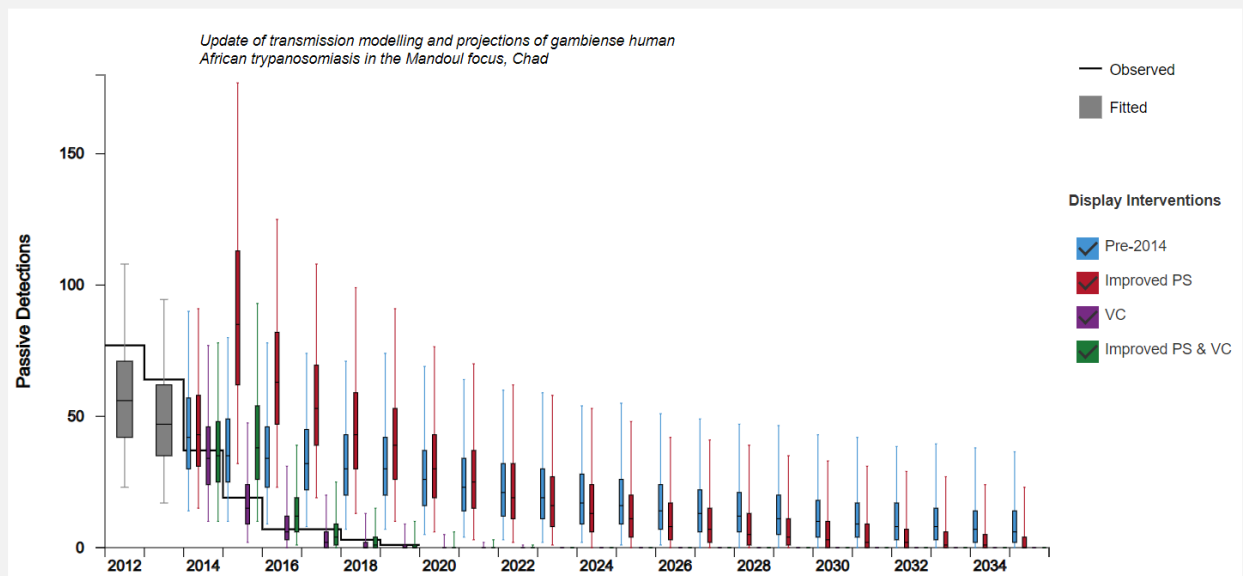
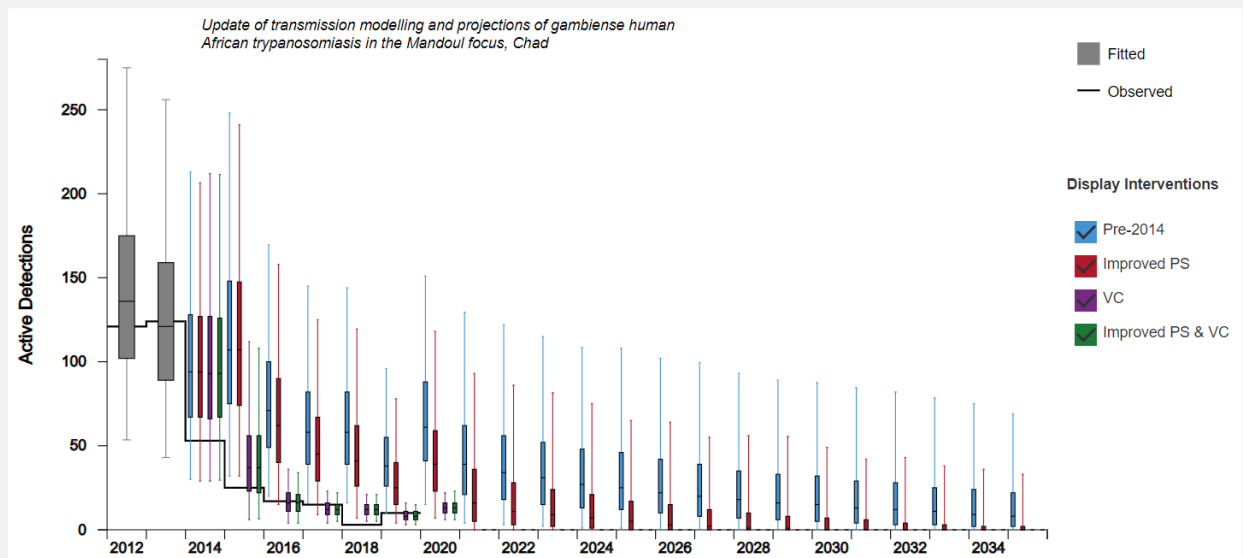


Illustration 5: Example results –new infections under four retrospective intervention strategies (2012 to 2035)

