# FleXScan User Guide

for version 3.0

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User Guide version 3.0e

#### Introduction

The FleXScan software has been developed to analyze spatial count data using the flexible spatial scan statistic developed by Tango and Takahashi (2005) and Kulldorff's circular spatial scan statistic (1997). The current version includes a spatial scan statistic tic with a restricted likelihood ratio proposed by Tango (2008). It is similar to the SaTScan software (2008) developed by Kulldorff together with Information Management System Inc., but the current version of FleXScan is still restricted to spatial analyses.

Kulldorff's circular spatial scan statistic uses a "circular window" with variable size to define the potential cluster area and so it is difficult to correctly detect noncircular clusters, such as those along a river. In real applications, most geographical areas are noncircular. Tango and Takahashi's flexible spatial scan statistic, therefore, is designed so that the detected cluster is allowed to be flexible in shape, while at the same time the cluster is confined within relatively small neighborhoods of each region. It should be noted that both of these scan statistics are based on maximizing the likelihood ratio. Tango (2000) showed an interesting example in which Kulldorff's circular spatial scan statistic detected an unrealistically large *most likely cluster* (MLC) consisting of 70 regions. This was much larger than expected from an observed disease map, and was due to absorption of neighboring regions with a non-elevated risk of disease occurrence in his simulated data. Furthermore, Tango and Takahashi have shown examples in which Duczmal and Assunção's non-circular scan statistic (2004) detected quite large and peculiar shaped MLC that had the largest likelihood ratio among the MLCs identified by the three different spatial scan statistics: Kulldorff's, Duczmal and Assunção's, and Tango and Takahashi's. These results cast doubt on the validity of model selection based on maximizing the likelihood ratio. Tango (2008) proposed a new spatial scan statistic free from this undesirable property by modifying the likelihood ratio so that it scans only the regions with elevated risk at the "region's significance level of alpha," specified by the user.

The FleXScan software is a free software program designed for any of the following interrelated purposes:

- 1) To evaluate reported spatial disease clusters, to see if they are statistically significant.
- 2) To test whether a disease is randomly distributed over space.

3) To perform geographical surveillance of disease, to detect areas of significantly high rates.

FleXScan uses the Poisson model, where the number of events in an area is Poisson distributed according to a known underlying population at risk. This version can also analyze data under the Binomial model. The data may be either aggregated at the census tract, zip-code, county or other geographical level. FleXScan can adjust for the underlying inhomogeneity of a background population and for any number of categorical covariates provided by user.

#### References

- Tango T and Takahashi K. (2005). A flexibly shaped spatial scan statistic for detecting clusters. International Journal of Health Geographics 2005, 4:11. [Open Access] <u>http://www.ij-healthgeographics.com/</u>
- Kulldorff M. (1997). A spatial scan statistic. Communications in Statistics: Theory and Methods, **26**:1481-1496.
- Tango T. (2008). A spatial scan statistic with a restricted likelihood ratio. Japanese Journal of Biometrics **29**:75-95.
- Kulldorff M and Information Management Services, Inc (2008). SaTScan<sup>TM</sup> v7.0: Software for the spatial and space-time scan statistics. <u>http://www.satscan.org/</u>
- Duczmal L and Assunção R. (2004). A simulated annealing strategy for the detection of arbitrarily shaped spatial clusters. Computational Statistics & Data Analysis, 45, 269-286.
- Tango T. (2000). A test for spatial disease clustering adjusted for multiple testing. Statistics in Medicine, **19**, 191-204.

# **Downloading and Installation**

To install FleXScan, go to the Department of Technology Assessment and Biostatistics, National Institute of Public Health Web site at :

http://www.niph.go.jp/soshiki/gijutsu/index\_e.html

and select the download link. After downloading the FleXScan installation executable to your PC, click on its icon and install the software by following the step-wise instructions.

System requirements:

- Intel® Pentium® processor III or higher
- Microsoft® Windows® 2000, Windows XP or Windows VISTA
- 256MB of RAM (recommended)

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Coordinate File:	
	Edit
Coordinates: 🕝 Latitude/Longitude 🔿 Cartesian	Radius of Earth: 6370 km
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# Test Run

Before using your own data, we recommend trying the sample data set provided with the software. To perform a test run:

- 1. Click on the FleXScan application icon.
- 2. Select the following files in the sample folder (C:/Program Files/FleXScan/sample, default) as input data.
  - ① Coordinate File: "saitama-e.coo"
  - ② Check 'latitude/Longitude' in the Coordinates
  - ③ Matrix Definition File: "saitama-e.mtr"
  - ④ Case File: "heart-M.cas"
- 3. Click on 'Set default name' or input any file name to output the results.

5 FleXScan	
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iles Analysis	
Input Coordinate File:	
C:¥Program Files¥FleXScanV3j¥sample¥saitama-e.coo	Edit
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C:¥Program Files¥FleXScanV3j¥sample¥saitama-e.mtr	Edit
	Edit
Case File ( observed # and expected # / population # ):	
C:¥Program Files¥FleXScanV3j¥sample¥heart-M.cas	Edit
Output	
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- 4. Next, click on 'Analysis' tab
  - ① Select "Poisson" in the 'Statistical model.'
  - 2 Select 'Original LLR' or 'LLR with Restriction' in the Statistic type, and 'Flexible' or 'Circular' in the 'Scanning method', respectively.

iles Analysis	Poisson C Binomial
Statistical model: Statistic type:	• Original LLR     C     LLR with Restriction     Alpha=     0.2     (default=0.2)
Scanning method: The Maximum Spat	Image: Flexible       Image: Circular         ial Cluster Size:       15         census areas (default=15, recommended<=20)
Random number: Monte Carlo replicat	Multinomial C Poisson

- 5. Click on 'Run' button (💈) or select 'Session' 'Run'.
- 6. After a while, the results file and the results map will appear.

Results:

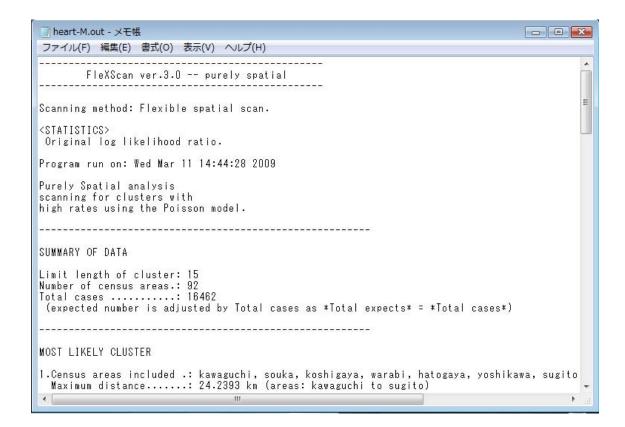
'MOST LIKELY CLUSTER'

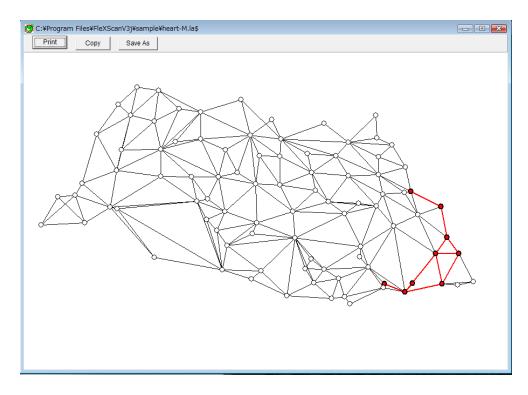
Census areas included. : kawaguchi, souka, koshigaya,...

are regions detected as the most likely cluster, and its significance is also given by

P-value .....: 0.001

Regardless of the p-value, the most likely cluster is shown in red on the map.





### Sample Data Sets and Data Input Form

To analyze your own data using FleXScan, you need to prepare at least three data files: 1) Coordinate File, 2) Matrix Definition File, and 3) Case File. The detailed structure of each file is explained in the following section showing sample data files for Saitama prefecture in Japan (automatically installed in your 'sample' folder).

① Coordinate File (coo)

The coordinate file provides the geographic coordinates for each area. Coordinates may be specified either using the standard 'Cartesian coordinates' system or in 'latitude and longitude.' 'Cartesian' is the regular planar x,y-coordinate system. Each line of the file represents an area name (or code) and its geographical location.

For the Cartesian coordinates system

Format : <Area name or code> <X-coordinate> <Y-coordinate>

For the Latitude and Longitude coordinates system

Format : <Area name or code> <Latitude> <Longitude>

Latitudes and longitudes should be entered as decimal numbers of degrees. You can convert latitudes and longitudes expressed in degrees, minutes, and seconds to decimal number of degrees by the following formula:

xx (degrees) yy (minutes) zz (seconds)  $\rightarrow$  xx + yy/60 + zz/3600 (degrees).

When coordinates are specified in latitudes and longitudes, FleXScan calculates the distance between two points on the surface of the spherical earth with a radius given in the 'Radius of Earth.'

File ( <u>F</u> )	(Scan¥sample¥ Edit (E)	saitama-e.co	0	
lo.	Area name	Latitude	Longitude	
	kawagoe		139.4891667	
	kumagaya		139.3919444	
:	kawaguchi		139.7272222	
	urawa	35.85833333		
i	oomiya		139.6319444	
			139.4588889	
	eyouda chichibu	35.98861111		
	tokorozawa	35.79638889		
	hannou	35.8525	139.3311111	
0	kazo		139.6052778	
1	honjyo	36.24055556		
2		36.03888889		
3	iwatsuki		139.7027778	
4	kasukabe		139.7558333	
5	sayama		139.4155556	
6	hanyu		139.5519444	
7	kounosu		139.5255556	
8	fukaya	36.19444444	139.2847222	
9	ageo	35.97416667	139.5966667	
0	yono	35.88055556	139.6291667	
1	souka	35.82222222	139.8086111	
2	koshigaya	35.88777778	139.7941667	
3	warabi	35.8225	139.6827778	
4	toda	35.81444444	139.6811111	
5	iruma	35.83277778	139.3944444	
6	hatogaya	35.82361111	139.7444444	
7	asaka	35.79416667	139.5969444	
8	siki	35.83333333	139.5836111	
9	wakou	35.77833333	139.6088889	
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2 Matrix Definition File (mtr)

➢ Format : <Area name or code> <Area 1> <Area 2>...

The first column of each line is the area name, which must be identical to that in Coordinate File. The following columns specify the area name(s) that are adjacent to (i.e., border on) the area described in the first column. For example, kawagoe, sayama, iruma, niiza, and miyoshi areas are adjacent to tokorozawa (see row No.8 of the figure below).

When Area1 is adjacent to Area3 and Area5, the mtr file should be:

Area1 Area3 Area5 Area2 .... Area3 Area1 ... ...

Note that "Area1" also appears in the line of "Area3" in this case (and vice versa), and the matrix must be symmetrical, otherwise an error occurs. The 'Check symmetry' tool in the File menu is available to check the symmetry of the matrix.

No.	Area name	Connected	Connected	Connected	Connected	Connected	Connected	Connected	
1	kawagoe	oomiya	tokorozawa	sayama	ageo	fujimi	kamifukuoka	sakado	
2	kumagaya	gyouda	higashimats	fukaya	fukiage	namekawa	oosoto	kounan	
3	kawaguchi	urawa	iwatsuki	souka	koshigaya	warabi	toda	hatogaya	
4	urawa	kawaguchi	oomiya	iwatsuki	yono	warabi	toda	asaka	
5	oomiya	kawagoe	urawa	iwatsuki	ageo	yono	fujimi	hasuda	
ŝ	gyouda	kumagaya	kazo	hanyu	kounosu	fukiage	menuma	kisai	
7	chichibu	naguri	tokigawa	yokose	minano	yoshida	okano	arakawa	1
8	tokorozawa	kawagoe	sayama	iruma	niiza	miyoshi			
9	hannou	sayama	iruma	hidaka	moroyama	ogose	naguri	tokigawa	1
10	kazo	gyouda	hanyu	kuki	kisai	kitakawabe	ootone	kurihashi	
11	honjyo	fukaya	misato-mac	kodama	kamisato	okabe	1		
12	higashimats	kumagaya	sakado	namekawa	arashiyama	kawashima	yoshimi	hatoyama	
13	iwatsuki	kawaguchi	urawa	oomiya	kasukabe	koshigaya	hasuda	shiraoka	
14	kasukabe	iwatsuki	koshigaya	miyashiro	shiraoka	sugito	matsubushi	syouwa	
15	sayama	kawagoe	tokorozawa	hannou	iruma	hidaka			
16	hanyu	gyouda	kazo						
17	kounosu	gyouda	okegawa	kitamoto	fukiage	yoshimi	kisai	kawazato	
18	fukaya	kumagaya	honjyo	menuma	okabe	kawamoto	hanazono	yorii	
19	ageo	kawagoe	oomiya	okegawa	hasuda	ina	kawashima		
20	yono	urawa	oomiya						
21	souka	kawaguchi	koshigaya	yashio	misato-shi	yoshikawa	1		
22	koshigaya	kawaguchi	iwatsuki	kasukabe	souka	yoshikawa	matsubushi	1	
23	warabi	kawaguchi	urawa	toda	- 1995 (1995 505)				

#### ③ Case File (cas)

The frequency of disease in each area is described in Case File. The current version of FleXScan can analyze two types of data.

- 1 'observed number' and 'expected number,'
- ➢ Format: <Area name or code> <Observed no.> <Expected no.>

For this data, 'Poisson model' should be selected in the 'Statistical model' for the analysis.

- ② 'observed number' and 'population,'
- Format: <Area name or code> <Observed no.> <Population >

For this data, 'Binomial model' should be selected in the 'Statistical model' for the analysis.

The first column of each line is the area name, which must be identical to that in the Coordinate File. The second column is the observed number of diseases, and the third column is the expected number of diseases under the null hypothesis, or the background population at risk in each area.

When you use the Poisson model, you need to calculate the expected number by yourself, for example, in the same manner as standardized mortality ratio (SMR). In kawagoe (see No.1 row of the figure) there were 705 deaths and the age-standardized expected number of deaths was 719.4 (i.e., SMR = 719.4 / 705 = 1.02).

But, if you do not need standardization (e.g., you are interested in the crude death rate), you can analyze the case file of 'observed number' and 'population' using the 'Poisson model.'

No:	Area name	Observed	Expected
1	kawagoe	705	719.3877551
2	kumagaya	451	389.4645941
3	kawaguchi	1089	932.3630137
4	urawa	1002	1000
5	oomiya	1016	1048.503612
6	gyouda	277	234.5469941
7	chichibu	216	194.9458484
8	tokorozawa	678	690.4276986
9	hannou	256	218.0579216
10	kazo	205	168.8632619
11	honjyo	185	167.2694394
12	higashimats	241	205.4560955
13	iwatsuki	248	254.3589744
14	kasukabe	424	386.8613139
15	sayama	338	348.0947477
16	hanyu	233	161.8055556
17	kounosu	185	175.1893939
18	fukaya	335	263.1578947
19	ageo	426	431.1740891
20	yono	225	190.5165114
21	souka	528	412.8225176
22	koshigaya	648	568.4210526
23	warabi	230	181.2450749
24	toda	183	177.6699029

#### Important Note:

- All area names or codes and their order must be identical among 'Coordinate File,' 'Matrix Definition File,' 'Case File,' and 'Population File' (if necessary).
- The area name or code cannot include a space character. Use an under score or a hyphen in stead of a space character.

Good	10001
Good	New_York
N.G	New York

## Editing your data set

The data files 1) to 3) can be edited using FleXScan data editor. Enter the file name and click the 'Edit' button to execute the data editor. You can copy and paste your data from other software such as MS-Excel. It may be convenient to input your data on MS-Excel and copy and paste it to the FleXScan data editor.

1) Editing Coordinate File (coo).

- > Input the area name and its latitude and longitude in each column.
- > If x and y-coordinates are used, select 'Cartesian' on the 'Files' tab panel.
- > 'Save & return' to finish editing data.

2) Editing Matrix Definition File (mtr).

- Coordinate File must be made before starting to edit Matrix Definition File.
- Input the area name, which must be identical to that of Coordinate File. It will be convenient to Copy & Paste all the area names from Coordinate File to Matrix Definition File.
- Select an area name and execute 'Edit Area List,' then a list of area names will appear in the order of distance from the selected area. Check the check-box of areas that are adjacent to the selected area. By clicking the 'OK' button, the checked areas will be automatically added to the 'Adjacent' columns (see the figure below).

File ( <u>F</u> )	Edit ( <u>E</u> )								
No.	Area name	Connected	Connected	Connected	Connected	Connected	Connected	Connected	Con
3	tokorozawa	kawagoe	sayama	iruma 🛛	niiza	miyoshi	-	-	
)	h	3			moroyama	ogose	naguri	tokigawa	yokc
0	k 🖾 toko	rozawa	_		kisai	kitakawabe	ootone	kurihashi	wasl
1	h Check th		COK C	ancel	kamisato	okabe			
2	h	ed areas.	S	va	arashiyama	kawashima	yoshimi	hatoyama	oosc
3	iv Invos iv Invos	:nı		^	kasukabe	koshigaya	hasuda	shiraoka	
4	k 🗹 sayam			Þ	shiraoka	sugito	matsubushi	syouwa	
5	s ✔iruma s ✔niiza				iruma	hidaka			
6	h 🗌 fujimi			=					
7	k kamifu siki	ukuoka			fukiage	yoshimi	kisai	kawazato	syou
8	ft 🗌 asaka				okabe	kawamoto	hanazono	yorii	
9	a 🗌 wakou				hasuda	ina	kawashima		
0	y hanno								
1	s tsuru hidaka	gashima			misato-shi	yoshikawa			
2	k vono	3		е	souka	yoshikawa	matsubushi		

The symmetry can be tested by executing 'File – Check symmetry.' If the information is not symmetrical (e.g., Area3 is selected as an adjacent area to Area1, but Area1 is not selected as an adjacent area to Area3), an error message will appear.

Supplementary technical information:

The Matrix Definition File is designed in a user-friendly format. FleXScan will automatically convert it to a Connection Information Matrix File (mt0 file) and utilize it for the calculation. You can see the format of the mt0 file using a text editor. Conversely, if you already have the area connection data in the format of an mt0 file, it can be converted to an mtr file by executing 'Tool – Matrix-file converter.'

3) Editing Case File (cas).

- > Coordinate File must be made before starting to edit Case File.
- > Case File can be edited in the same manner as Coordinate File.

## **Parameters**

You can change several parameters for the analysis on the 'Analysis' tab panel.

- Statistical model
  - ① Poisson: for the data of the 'observed number' and the 'expected number.'
  - ② Binomial: for the data of the 'observed number' and the 'population.'
- Statistic type
  - ① Original LLR:

The likelihood ratio statistic by Kulldorff, which has been used in the previous FleXScan version 1 and version 2.

② LLR with Restriction:

The restricted likelihood ratio statistic by Tango, with a preset parameter for restriction 'Alpha' (default is 0.2). This statistic avoids detecting undesirably large clusters, and improves calculation time. See the reference article for details.

- Scanning method
  - ① Flexible : flexible scan statistic by Tango and Takahashi
  - ② Circular : circular scan statistic by Kulldorff
- The Maximum Spatial Cluster Size

The number of maximum spatial cluster size to scan. See the reference article for details.

• Random number:

The type of random number for Monte Carlo simulation

- ① Multinomial: Total number of cases in whole area is fixed. It can be chosen in either 'Poisson' or 'Binomial' model.
- ② Poisson: Total number of cases is not fixed, and it can be chosen in 'Poisson model'
- ③ Binomial: Total number of cases is not fixed, and it can be chosen in 'Binomial model'

• Monte Carlo replications

The number of Monte Carlo replications to calculate a p-value for statistical test. For example, if this number is set to 999, the p-value is calculated from the simulated 999 plus 1 observed log-likelihood ratio values (999+1=1000 in total).

• Seed of Random Number

The seed for generating random numbers in the Monte Carlo simulation.

😏 FleXScan		
File (F) Session (S)	Tool (T) Help (H)	
Statistical model: Statistic type:	Poisson C Binomial     Original LLR C LLR with Restriction Alpha= 0.2	(default=0.2)
Scanning method: The Maximum Spati	<ul> <li>☞ Flexible</li></ul>	
Random number: Monte Carlo replicat	<ul> <li>Multinomial C Poisson</li> <li>ons (default=999): 999 Seed of Random Number: 45861</li> </ul>	11

Also, on the 'Files' tab panel,

• Coordinates

The type of coordinates used by the coordinates file.

• Radius of Earth

Radius of Earth to calculate a distance between two sets of latitude and longitude. It is approximately 6370 km in Japan.

🗭 FleXScan File (F) Session (S) Tool (T) Help (H)	_ • •
Files Analysis	
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Matrix Definition File:	
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Case File ( observed # and expected # / population # ):	
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## Software Licenses

- The FleXScan software may be used freely, with proper references to both the software and the statistical methods papers. The suggested citations are:
  - Tango T. and Takahashi K. A flexibly shaped spatial scan statistic for detecting clusters. International Journal of Health Geographics 2005, 4:11.
  - Takahashi K, Yokoyama T and Tango T. FleXScan: Software for the Flexible Scan Statistic. National Institute of Public Health, Japan, 2005 - 2009.

Also for the spatial scan statistic with a restricted likelihood ratio,

Tango T. A spatial scan statistic with a restricted likelihood ratio. Japanese Journal of Biometrics 2008, 29:75-95.

# **Copyrights**

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http://www.niph.go.jp/soshiki/gijutsu/index\_e.html