



# embadpixfind

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## Abstract

Find EPIC-MOS bad pixels for one node of one EPIC-MOS CCD.

## 1 Instruments/Modes

Instrument	Mode
EPIC MOS	IMAGING

## 2 Use

pipeline processing	yes
interactive analysis	yes

## 3 Description

**embadpixfind** aims at finding bad pixels in an EPIC-MOS image in a completely automatic way, by taking advantage of the broad PSF in pixel units, which makes it impossible to mistake a source for a bad pixel.

### 3.1 Preparatory work

Call **emeventsproj** with `rejectbadevents=Y` to project the events file to an image. An image built by **evselect** would be accepted as well, but would not include the secondary pixels of multiple events (this may miss weak bad pixels next to a strong one).

For optimal electronic noise rejection, the events file should be built by successive calls to **emevents** (`analysepatterns=N flagbadpixels=N splitdiagonals=N randomizeposition=N` allow it to run much faster) and **emenergy** (`correctcti=N correctgain=N randomizeenergy=N` allow it to run much faster).

For calibration purposes, it is possible to improve the statistics (and the sensitivity to weak bright pixels or dark pixels) by stacking many images output of **emeventsproj** on top of one another before calling **embadpixfind**.



### 3.2 Basic algorithm

**embadpixfind** estimates the local statistical average  $\mu$  in a running window around each pixel by taking the smallest of the average or the median + 1 (1 is added to take care of the case when the median is 0, the median allows to remove the effect of other bad pixels in the vicinity). Then it builds a significance map via the Li and Ma criterion (Li & Ma 1983, ApJ 272, 317):

$$S = \sqrt{2} \sqrt{N_{\text{on}} \ln \frac{N_{\text{on}}}{\mu_{\text{tot}}} + N_{\text{off}} \ln \frac{\mu}{\mu_{\text{tot}}}} \quad (1)$$

where  $N_{\text{on}}$  is the number of counts in the current pixel,  $N_{\text{off}} = N_{\text{pix}} \mu$  is the number of reference counts,  $N_{\text{pix}}$  is the number of pixels used to compute the local average ( $(2 \text{ halfwidth2d} + 1)^2 - 1$ , if none of the pixels in the window has been rejected already),  $N_{\text{tot}} = N_{\text{on}} + N_{\text{off}}$  is the total number of counts in the window, and  $\mu_{\text{tot}} = N_{\text{tot}} / (N_{\text{pix}} + 1)$  is the average number of counts per pixel in the window.

This significance map is then used to locate the most promising candidate bad pixels. They are examined in turn, in decreasing order. The exact probability that the current excess is a statistical anomaly of a flat distribution is computed from the cumulative binomial law:

$$P(k \geq N_{\text{on}}) = \sum_{k=N_{\text{on}}}^{N_{\text{tot}}} p_B(k, N_{\text{tot}}, q) = I_q(N_{\text{on}}, N_{\text{off}} + 1) \quad (2)$$

where  $q = 1 / (N_{\text{pix}} + 1)$  is the probability that a random count fall in the central pixel, and  $I_x(a, b)$  is the incomplete beta function. This is significantly different from the probability estimated from Eq.(1) for small numbers (Eq.3 gives a larger probability). If that probability is smaller than **probathreshold**, the pixel is flagged as bright, the average is recomputed around the bad pixel ignoring it, and the loop goes on. The loop stops when the next largest excess is smaller than the significance corresponding to **probathreshold**.

### 3.3 Columns and rows

If **findbadsegments=Y embadpixfind** looks for bright rows and columns too. This is done by projecting the image along rows and columns, and applying the same algorithm as above on the resulting 1-D vectors, except the width of the 1-D window ( $2 \text{ halfwidth1D} + 1$ ) is normally chosen larger than that of the 2-D window, to improve the background determination.

If a bright row or column is found, it is analysed to look for bright segments within. If it is found that the rest of the row/column is compatible (to 1% probability) with the neighbouring rows/columns, then only the bright segments are declared as bad. The minimum length of the bright segments is set such that one expects about 1 count in that length in the normal (not bright) parts.

### 3.4 Intrinsic dispersion

Sometimes the distribution of the number of counts in the window around the tested pixel does not follow the Poisson law at all, but is much broader. This is particularly true for columns at low energy, because of charge transfer efficiency variations from one column to the next. This effect is of course more obvious for large count rates, in particular within bright sources.

To avoid wrongly detecting columns as bad (either dark or bright), the observed dispersion in the distribution is used to compute the significance of an excess assuming a Gaussian distribution. What is actually measured is the average absolute deviation (this is more robust than the root mean square when a few



other bright pixels are present) divided by 0.8 (to recover the standard deviation when the distribution is normal). The average absolute deviation is estimated discarding (of course) the tested column, but also the highest one among others, to allow detecting pairs of bright columns. The true significance is taken to be the smallest of the Gaussian estimate and the Poisson one (from Eq.1). The cost of this security is to detect less easily groups of bad columns/rows, because the observed dispersion is large even when the tested column/row is discounted.

In addition, the `minratio` parameter avoids detecting bright pixels, rows or columns with too small contrast on observations with high statistics.

### 3.5 Dark features

If `finddead=Y`, `embadpixfind` looks for too dark pixels, rows and columns too. The Li and Ma formula (Eq.1) is not used for dark pixels. The Gaussian significance is computed in the same way (but on the negative side) and the cumulative binomial probability is computed from:

$$P(k \leq N_{\text{on}}) = \sum_{k=0}^{N_{\text{on}}} p_B(k, N_{\text{tot}}, q) = I_{1-q}(N_{\text{off}}, N_{\text{on}} + 1) \quad (3)$$

The statistics in a single observation is usually not enough to find any dark pixel, but dark rows or columns may be found. The `maxratio` parameter avoids detecting 'grey' pixels, rows or columns on observations with high statistics.

Whatever `findbright` and `finddead`, dark and bright columns and rows are always searched for together in order of decreasing significance (either positive or negative). This avoids finding spurious bright columns/rows next to very dark ones, and vice-versa.

If one of `findbright` or `finddead` is set to False, the corresponding bad pixels/columns/rows are not written to the output file.

### 3.6 Iteration

Each step in the detection process may affect the other steps. For example, detecting a bright row or column may facilitate detecting a moderately bright pixel next to it. For that reason, the whole process is iterated until nothing new is detected, or until the maximum number of iterations (set by the `niter` parameter) is reached.

### 3.7 Calibration access

If `usecal= Y`, the uplinked bright pixels, as well as known bright and dead pixels (within the current window), are read from the CAL and those pixels are ignored in computing the local median and the threshold. Columns switched off by large offsets are also ignored and considered similarly to uplinked bright pixels.

Optionally (`includedeadpixels` parameter) one may include the dead pixels from the CAL in the output list. It is also possible (`ignoreccfbright` parameter) to ignore the bright pixels declared in the CCF and redetect them from the data. This does not apply to uplinked and dead pixels which appear dark in the data and cannot be detected in a single observation. If `embadpixfind` was called with `ignoreccfbright=Y` `includedeadpixels=Y`, `badpix` may be called with `getnewbadpix=Y` `getotherbadpix=N`



to keep only the bright pixels active in the current exposure while preserving the information about dead pixels.

With `usecal=N`, the `embadpixfind` algorithm is not XMM specific at all and works on any image where the normal structure size is larger than 5 pixels.

### 3.8 Incremental search

The default mode is to ignore the bad pixels file on input. If `incremental=Y`, `embadpixfind` reads the bad pixels file on input, and ignores the pixels mentioned there in the search. On output the bad pixels file contains both the original bad pixels and the newly found ones.

## 4 Parameters

This section documents the parameters recognized by this task (if any).

Parameter	Mand	Type	Default	Constraints
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<code>evimageset</code>	yes	dataset	' '	none
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Input image file (from `emeventsproj`)

<code>badpixset</code>	no	dataset	<code>badpix.out</code>	none
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Output bad pixels file

<code>incremental</code>	no	boolean	no	yes/no
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Add newly found bad pixels to the bad pixels file contents

<code>probathreshold</code>	no	real	$1.10^{-6}$	$> 0, < 1.10^{-3}$
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False detection probability per pixel

<code>halfwidth2d</code>	no	integer	2	$> 0$
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Half width for 2D searches (images)

<code>findbadsegments</code>	no	boolean	yes	yes/no
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Look for bad segments of rows or columns as well

<code>halfwidth1d</code>	no	integer	3	$> 0$
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Half width for 1D searches (columns/rows)

<code>findbright</code>	no	boolean	yes	yes/no
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Look for too bright pixels, rows and columns

<code>minratio</code>	no	real	1.5	$> 1$
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Minimum ratio to neighbours for bright features (when `findbright=Y`)

<code>finddead</code>	no	boolean	yes	yes/no
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Look for too dark pixels, rows and columns

<code>maxratio</code>	no	real	0.5	$> 0, < 1$
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Maximum ratio to neighbours for dark features (when `finddead=Y`)



<b>niter</b>	no	integer	10	> 0
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Maximum number of iterations of the full detection process

<b>usecal</b>	no	boolean	yes	yes/no
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Get uplinked and dead pixels from the CAL

<b>includedeadpixels</b>	no	boolean	no	yes/no
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Include dead pixels from CAL in output list

<b>ignoreccfbright</b>	no	boolean	no	yes/no
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Ignore the bright pixels declared in the CAL (except uplinked)

## 5 Errors

This section documents warnings and errors generated by this task (if any). Note that warnings and errors can also be generated in the SAS infrastructure libraries, in which case they would not be documented here. Refer to the index of all errors and warnings available in the HTML version of the SAS documentation.

### **getParamValues03** (*error*)

keyword incompatibility between image and bad pixels files (**incremental=Y**)

### **getCalBadpix10** (*warning*)

bright pixel wrongly declared as uplinked in the CCF. That pixel is treated as not uplinked.

*corrective action:* inform SOC (this is a CCF error)

### **getCalBadpix11** (*warning*)

offsets are not the same size as the expected window.

*corrective action:* inform SOC (this is a CCF or coding error)

## 6 Input Files

1. Projected image file (from emeventsproj) as Integer\*4 array in PRIMARY. If they exist, the window keywords WINDOWX0, WINDOWDX, WINDOWY0 and WINDOWDY are read to avoid underestimating the median and average at the borders in window mode.
2. File with BADPIX extension (if **incremental=Y**) output of **badpixfind**, **embadpixfind** or **badpix**.

## 7 Output Files

1. Bad pixels file (for **badpix**) as BADPIX extension with RAWX, RAWY, TYPE, YEXTENT and BADFLAG Integer\*2 columns.



## 8 Algorithm

```
Read the parameters

Define goodPixel array, set to True

if incremental then
  Read the bad pixels already in file
  goodPixel(bad pixels) = False
endif

if usecal then
  Read the bad pixels and offsets in CAL
  goodPixel(uplinked and dead pixels) = False
  if not ignoreccfbright then goodPixel(bright pixels) = False
  if includedeadpixels then Write dead pixels to output list
endif

Read map = projected image

Iterate
  call findAllBad(map, goodPixel, bad)
until nothing new is found or niter is reached

Write bad pixels file

subroutine findAllBad(map, goodPixel, bad)

! Get local average around each pixel and estimated significance
  call avMedFilter(map, goodPixel, medsmooth, badtest)
! Look for bright pixels
  call findBadPix(map, goodPixel, medsmooth, badtest, False, bad)
  if findbadsegments then
! Look for bright/dark columns
    profil = sum(map,2)
    call findBad1D(profil, bad)
! Look for bright/dark rows
    profil = sum(map,1)
    call findBad1D(profil, bad)
    Update goodPixel and medsmooth/badtest around bad columns/rows
  endif
! Look for dark pixels
  call findBadPix(map, goodPixel, medsmooth, badtest, True, bad)

end subroutine findAllBad

subroutine avMedFilter(map, goodPixel, medsmooth, badtest)

  Loop over current pixel
  Consider all good pixels in window around current pixel
  Extract average (or median+1 if smaller) into medsmooth
```



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    Estimate dispersion from average absolute deviation divided by 0.8
    Estimate significance S1 of excess using Gaussian law
    If larger than 3, estimate significance S2 of excess using Li and Ma
    badtest = min(S1,S2)
endloop

end subroutine avMedFilter

subroutine findBadPix(map, goodPixel, medsmooth, badtest, negative, bad)

  Loop over current bad pixel
  Find maximum in badtest (minimum if negative) down to probathreshold
  Get probability of excess using binomial law
  if probability < probathreshold then
    add bad pixel to list
    Update goodPixel and medsmooth/badtest around maximum
  endif
endloop

end subroutine findBadPix

subroutine findBad1D(profil, psf1D, bad)

  Find bad columns in profil (same as findBadPix)
  Loop over bad columns
  Get expected distribution along column from its neighbours
  width = 1/(expected count rate per pixel)
  Get running integral over bins of width pixels
  While total(rest of column) larger (lower if negative) than expected
    Find maximum integral (minimum if negative)
    Remove segment of width pixels around it
  endwhile
endloop

end subroutine findBad1D
```

## 9 Comments

- The algorithm to get the local average could be improved over using the local median. An algorithm similar to that used in computing the offsets in **emdiag** should be considered.

## References