Subject: bimodal distribution

Posted by Giuseppe Papa on Fri, 19 Nov 2010 18:22:55 GMT

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I'm trying to simulate a bimodal "randomn" distribution with the 2 peaks having the sigmas of 0.3 each for example and mean around -1.2 and -0.7...

Do you have any tips?

I've tried adding these below, but it doesn't work..., I get a unimodal distribution...

 $met_1a = randomn(seed, 100000)*0.3-1.2$ 

 $met_1b = randomn(seed, 100000)*0.3-0.7$ 

I hope you can help me...

Cheers

Subject: Re: bimodal distribution

Posted by Dick Jackson on Tue, 23 Nov 2010 21:09:29 GMT

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Hi, ...

I think these do give a bimodal distribution, but they overlap a lot.

This might help, separating them by an extra two units:

met\_1a = randomn(seed,100000)\*0.3-1.2 -1

 $met_1b = randomn(seed, 100000)*0.3-0.7 +1$ 

plot,histogram([met\_1a,met\_1b],binsize=0.1),psym=10

Note: the X axis is off here, as you need to wrestle with Histogram() and Plot() to get them right, but I think this makes the point.

Cheers,

-Dick

Dick Jackson Software Consulting

http://www.d-jackson.com

## chiessantos wrote:

- > I'm trying to simulate a bimodal "randomn" distribution with the 2 peaks having the sigmas of 0.3 each for example and mean around -1.2 and -0.7...
- > Do you have any tips?

>

- > I've tried adding these below, but it doesn't work..., I get a unimodal distribution...
- > met 1a = randomn(seed,100000)\*0.3-1.2
- > met\_1b = randomn(seed,100000)\*0.3-0.7

>

> I hope you can help me...

```
> Cheers
```

Subject: Re: bimodal distribution Posted by David Fanning on Tue, 23 Nov 2010 21:25:57 GMT View Forum Message <> Reply to Message

## Dick Jackson writes:

- > I think these do give a bimodal distribution, but they overlap a lot.
- > This might help, separating them by an extra two units:

>

- > met\_1a = randomn(seed,100000)\*0.3-1.2 -1
- > met\_1b = randomn(seed,100000)\*0.3-0.7 +1
- > plot,histogram([met\_1a,met\_1b],binsize=0.1),psym=10

>

- > Note: the X axis is off here, as you need to wrestle with Histogram()
- > and Plot() to get them right, but I think this makes the point.

No wrestling needed with Histoplot. :-)

IDL> Histoplot, [met 1a, met 1b], Binsize=0.1

Cheers,

David

David Fanning, Ph.D.

Fanning Software Consulting, Inc.

Coyote's Guide to IDL Programming: http://www.dfanning.com/

Sepore ma de ni thui. ("Perhaps thou speakest truth.")