THIS IS MY .BIBI FILE

```
<?xml version="1.0" encoding="UTF-8"?>
<bibi xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance"
xmlns="http://schemas.humanbrainproject.eu/SP10/2014/BIBI"
xsi:schemaLocation="http://schemas.humanbrainproject.eu/SP10/2014/BIBI ../bibi_configuration.xsd">
<brainModel>
<file>brain_model/visual_segmentation.py</file>
</brainModel>
<bodyModel>icub_model/model.sdf</bodyModel>
<transferFunction xsi:type="PythonTransferFunction" src="TransferFunctions/grab_image.py"/>
<!-- <transferFunction xsi:type="PythonTransferFunction"
src="TransferFunctions/send_segmentation_signals.py"/> -->
<transferFunction xsi:type="PythonTransferFunction" src="TransferFunctions/plot_V2_activity.py"/>
<transferFunction xsi:type="PythonTransferFunction" src="TransferFunctions/plot_V4_activity.py"/>
<transferFunction xsi:type="PythonTransferFunction">



#### BELOW IS THE TRANSFER FUNCTION THAT DISPLAYS ALL NEURONS


#<![CDATA[
import hbp_nrp_cle.tf_framework as nrp
@nrp.NeuronMonitor(nrp.brain.V2Layer23, nrp.spike_recorder)
def boundary_segmentation_spike_monitor(t):
return True
#]]>
</transferFunction>



####



</bibi>
```