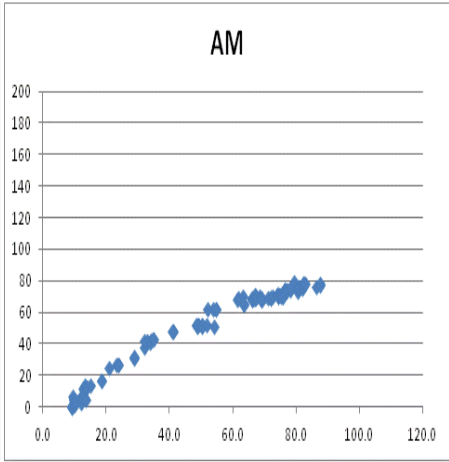
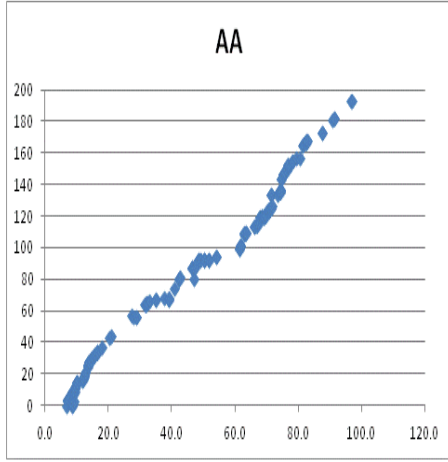
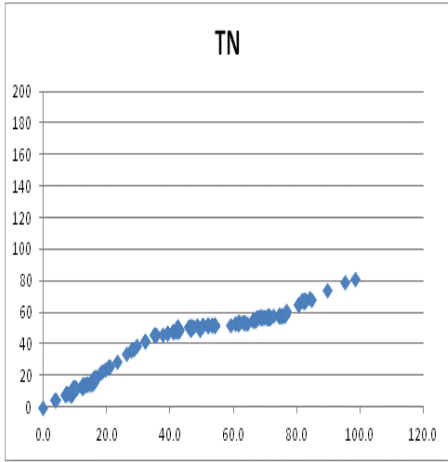
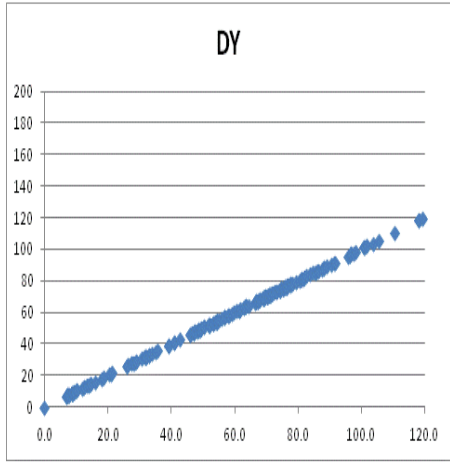
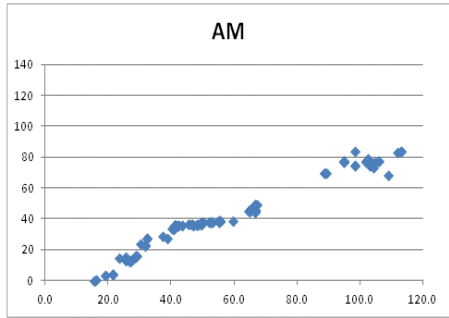
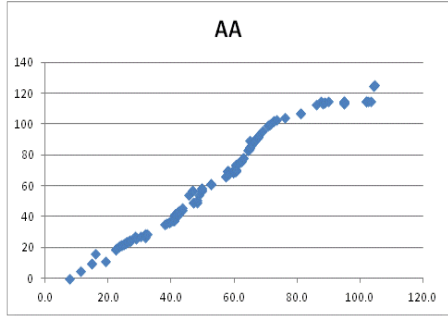
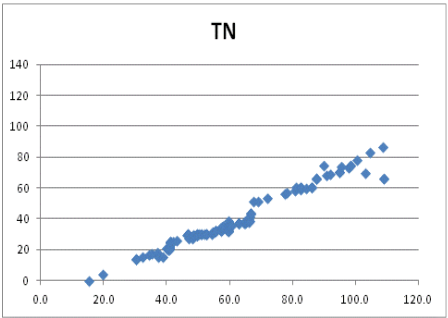
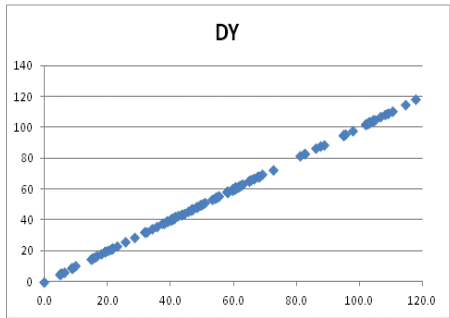


**Supplementary figure 2:** Dot-plots obtained for each linkage group between the integrated map and all four individual TNDH, DYDH, AADH and AMDH maps. The marker order on the vertical axis is from the four individual maps and the marker order on the horizontal axis is from the integrated map. Cumulative genetic distance in cM is indicated on each axis.

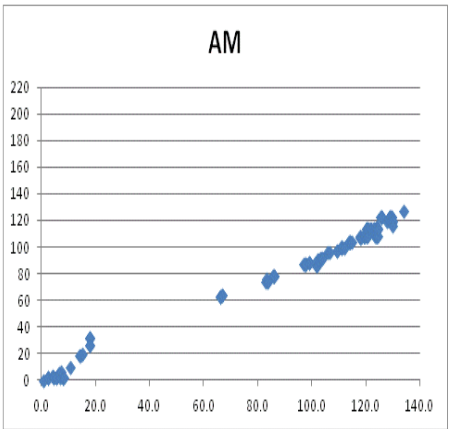
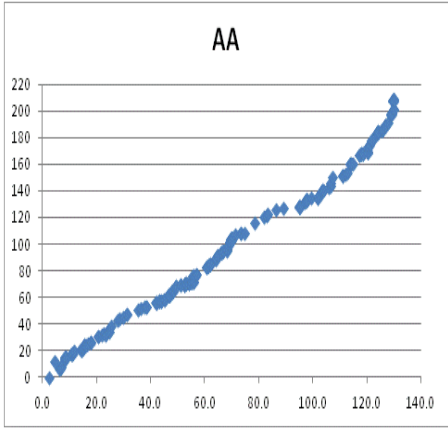
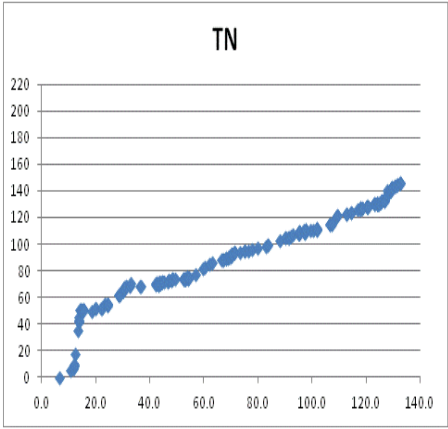
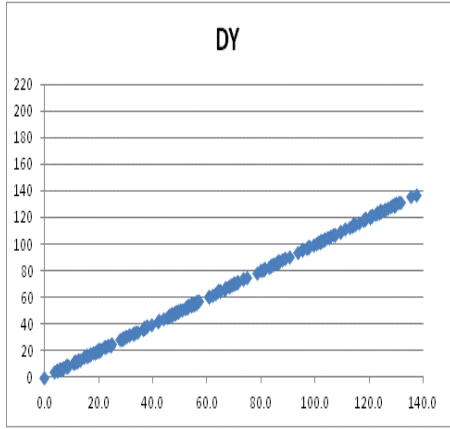
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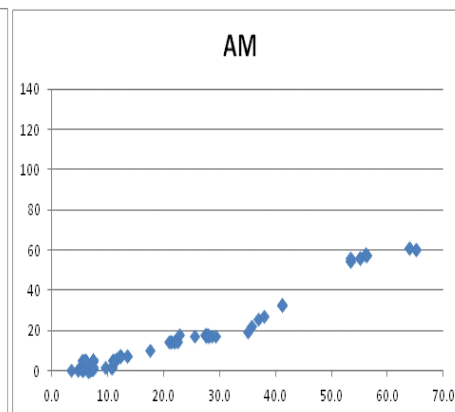
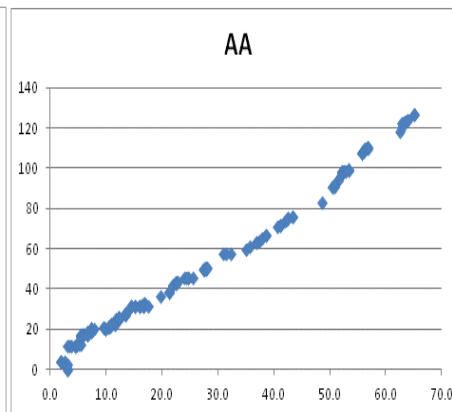
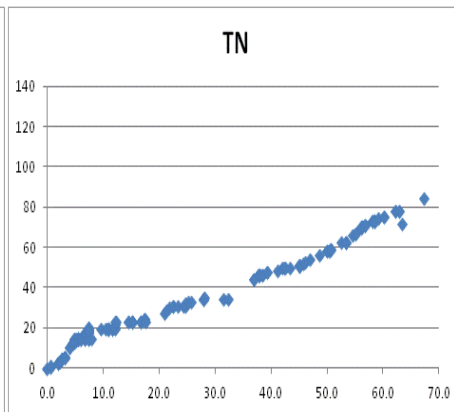
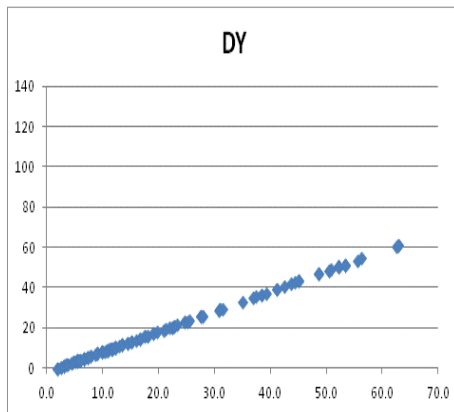
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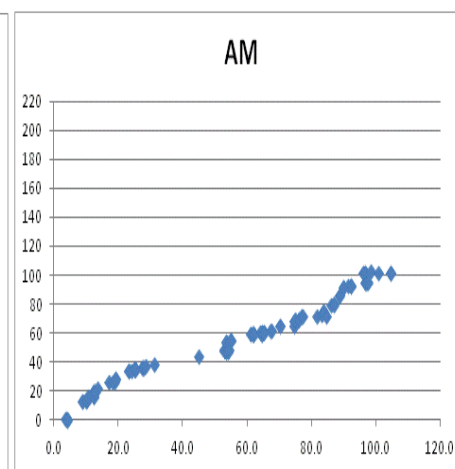
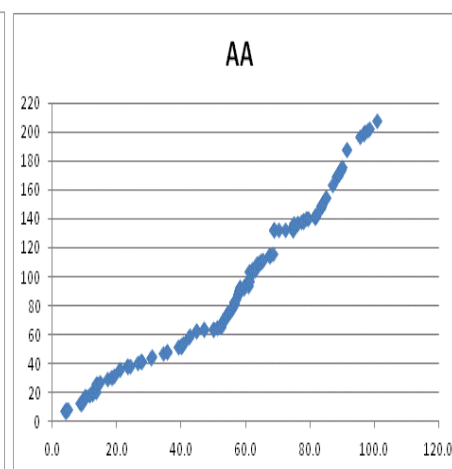
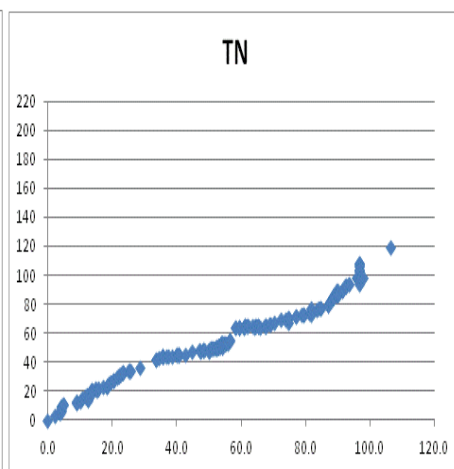
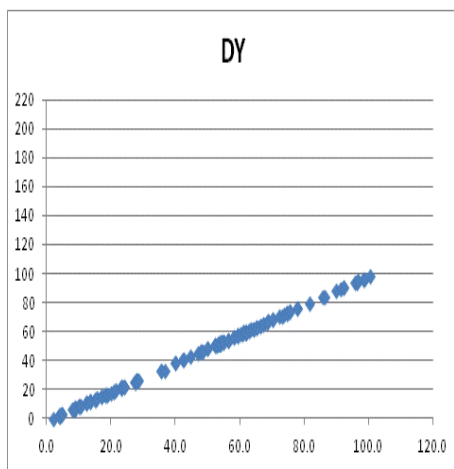
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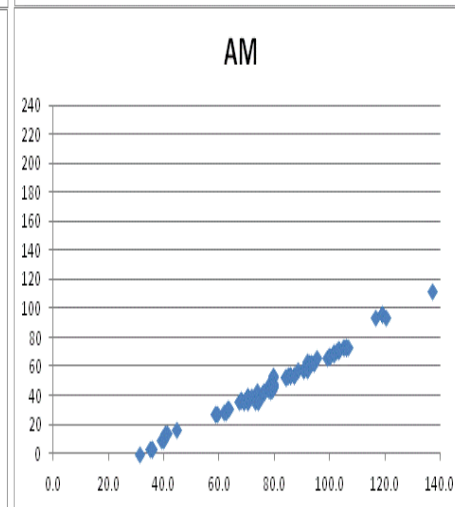
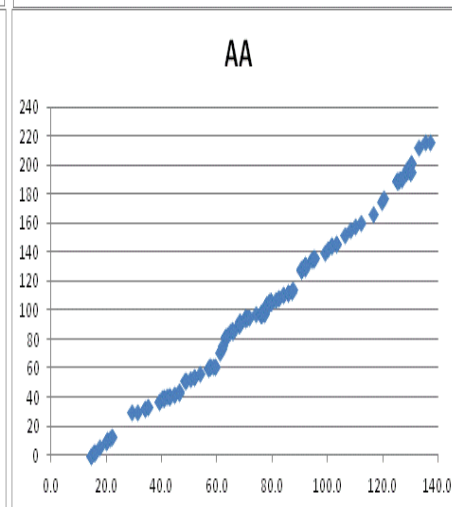
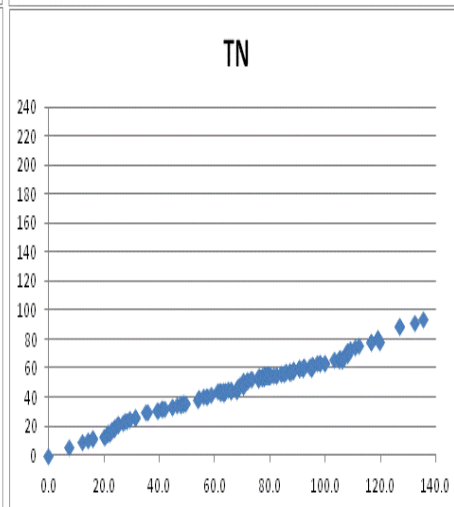
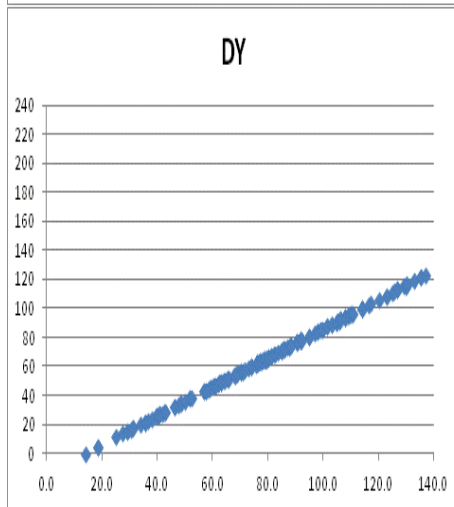
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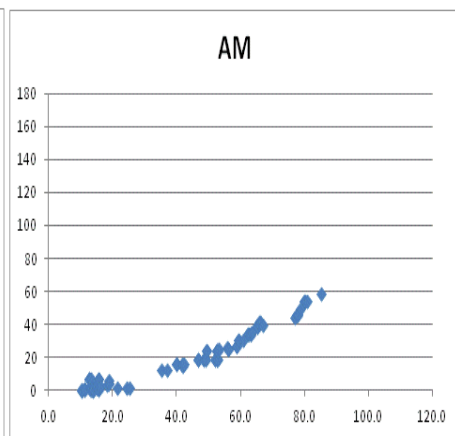
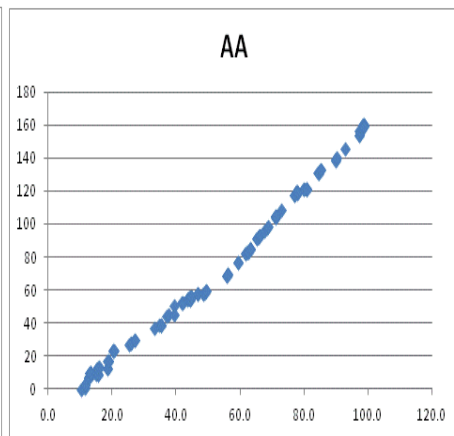
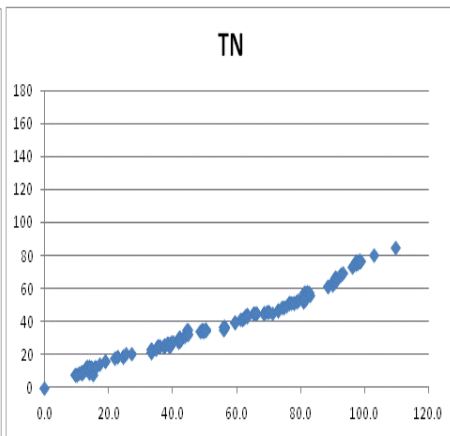
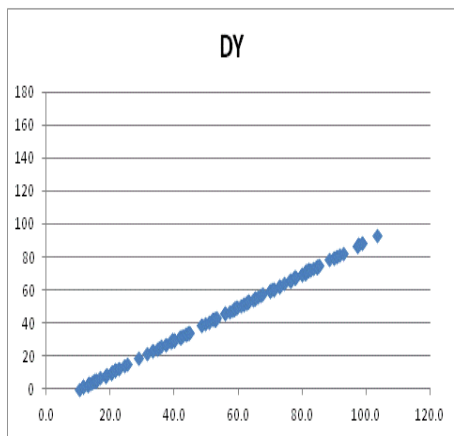
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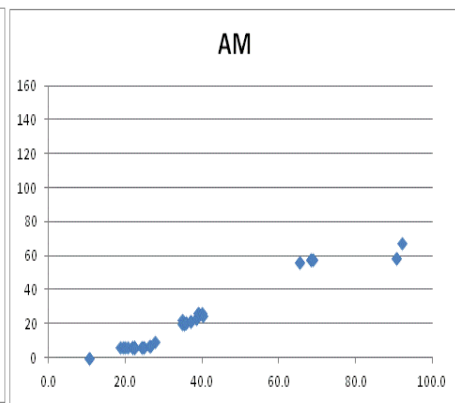
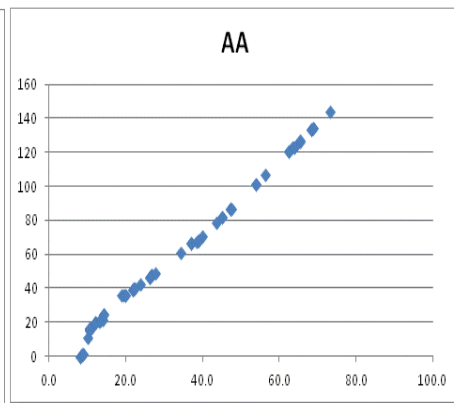
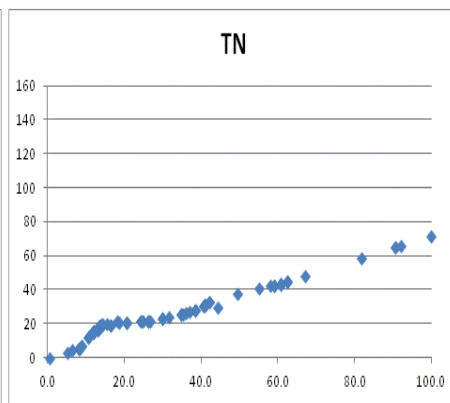
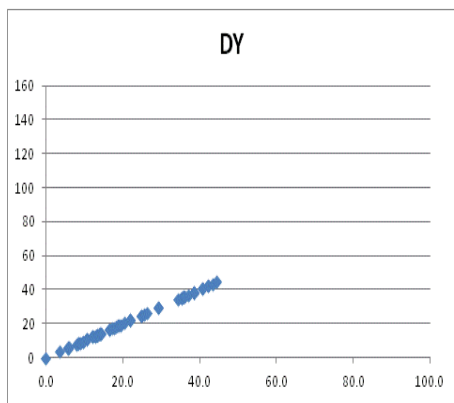
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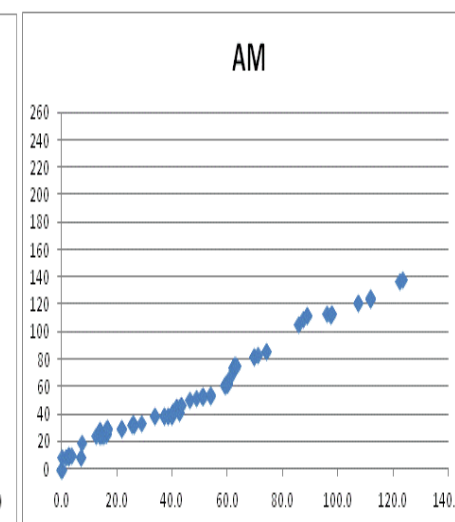
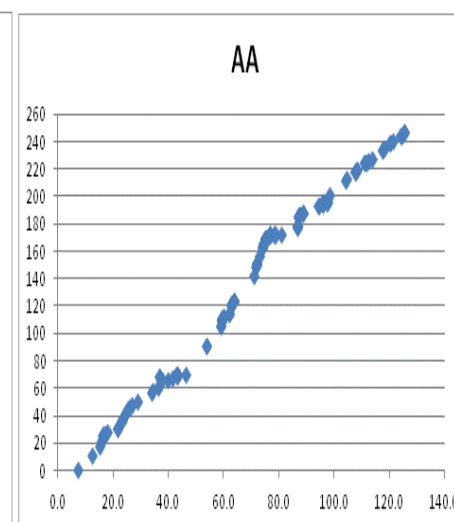
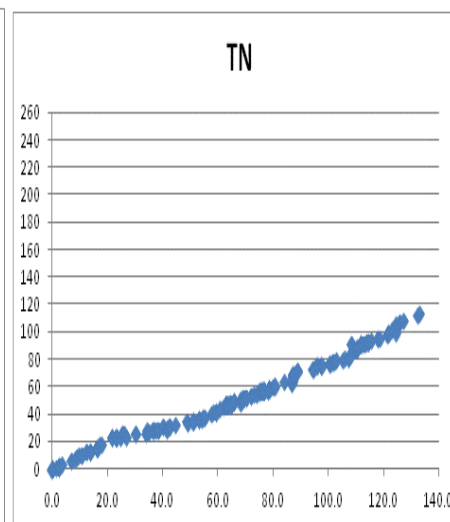
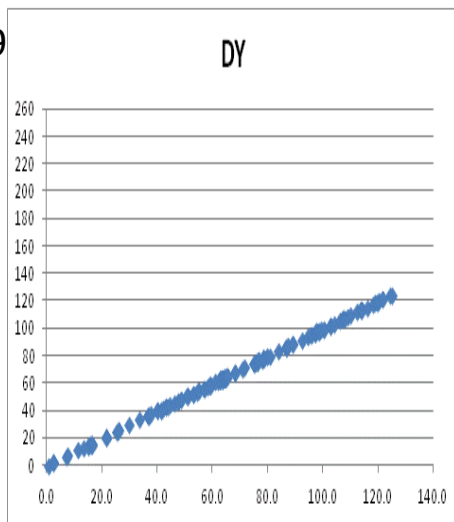
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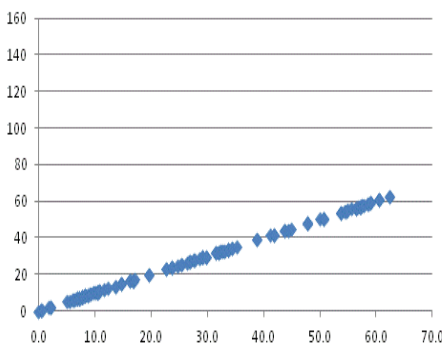


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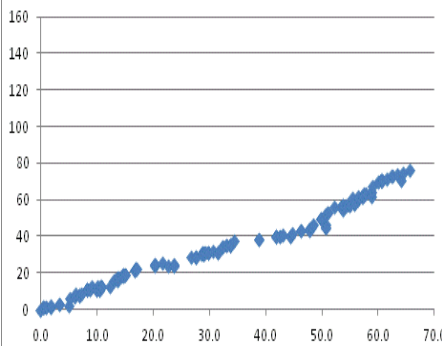


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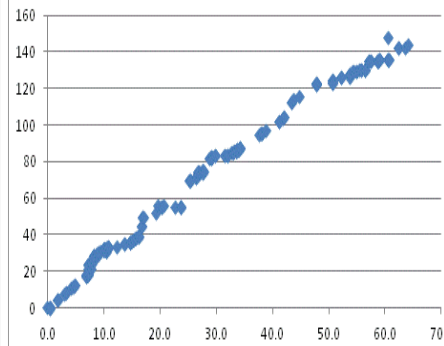
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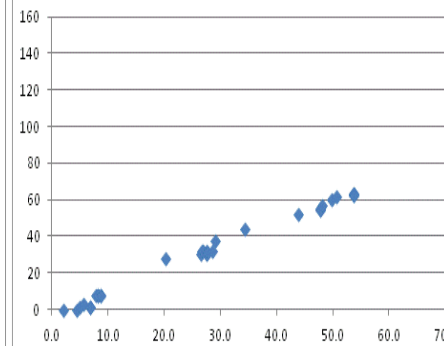
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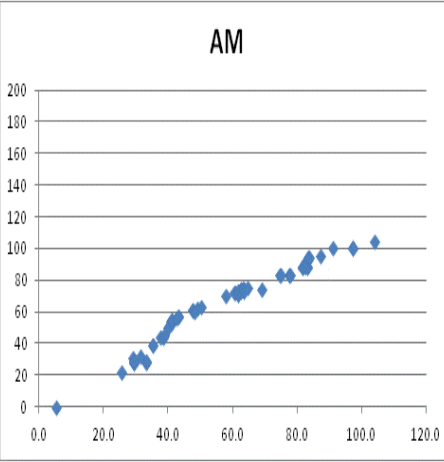
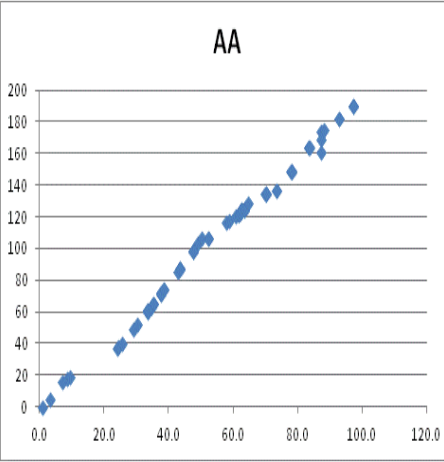
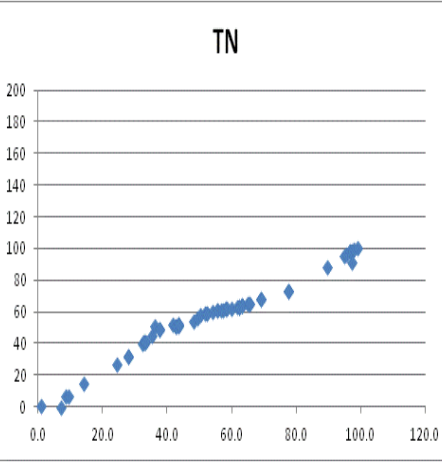
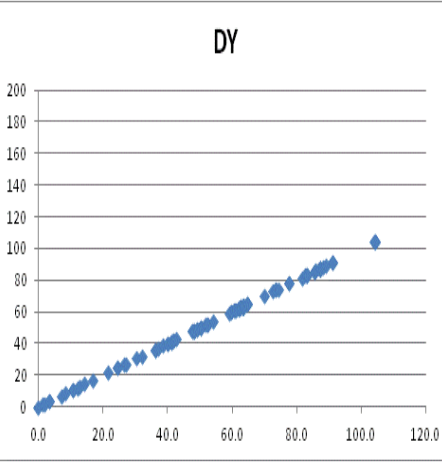
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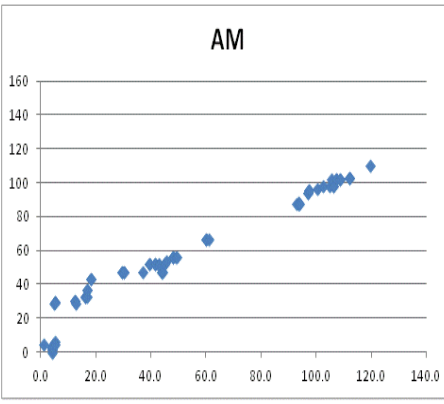
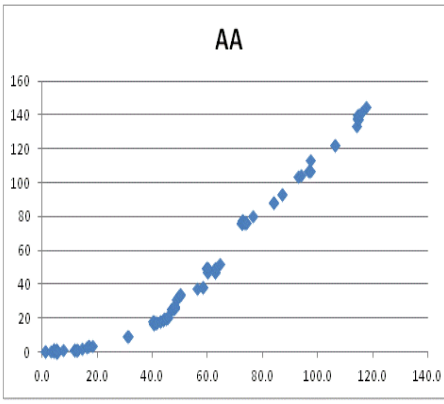
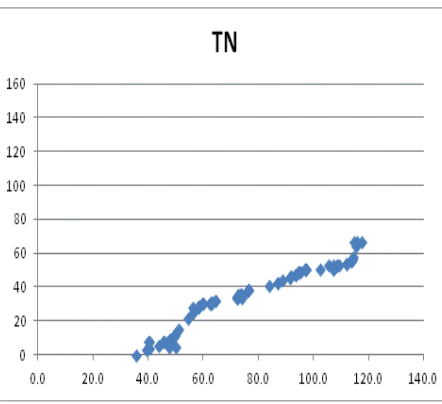
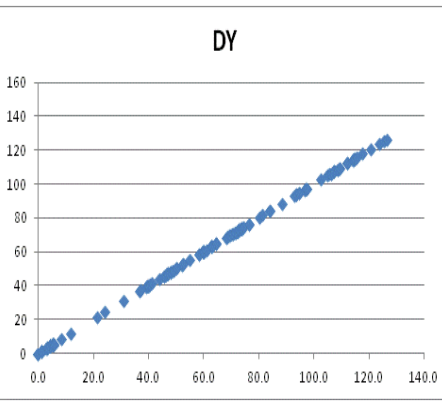
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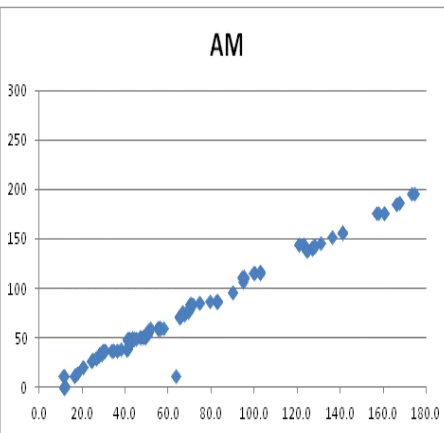
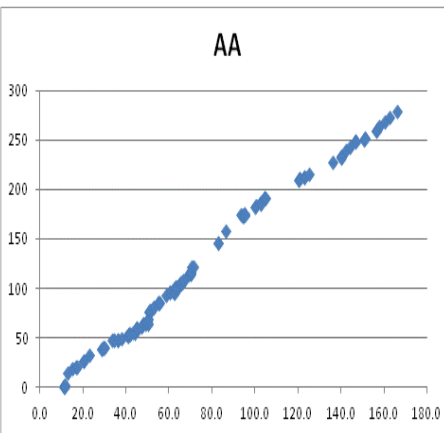
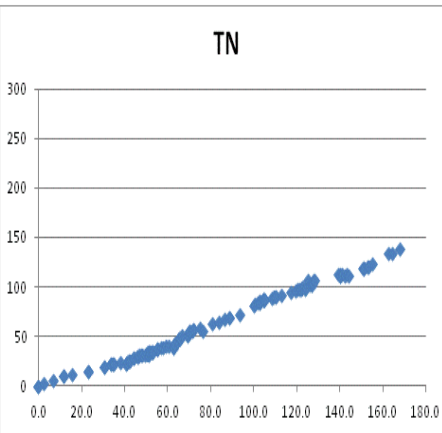
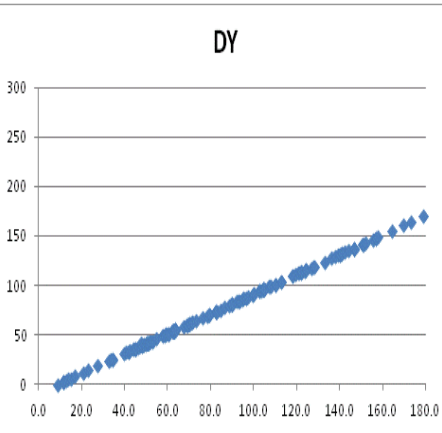
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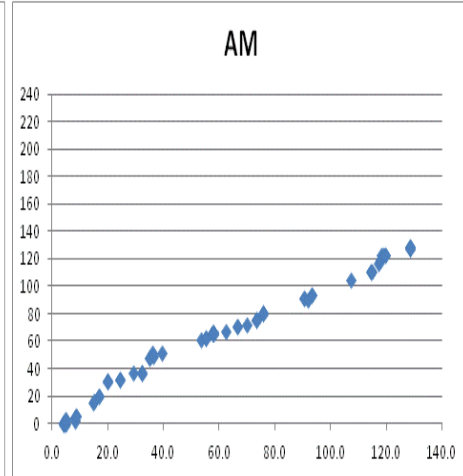
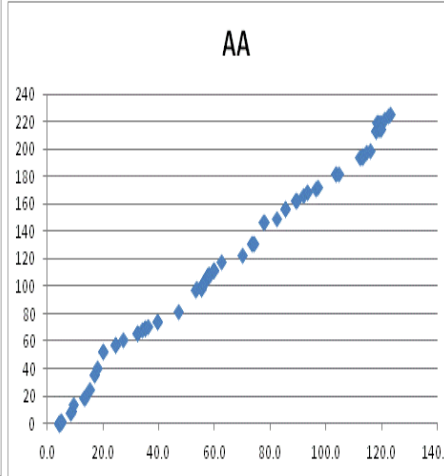
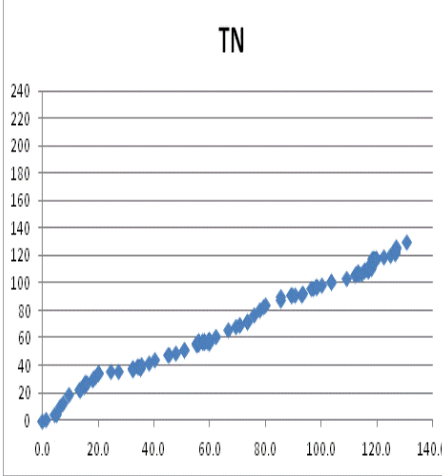
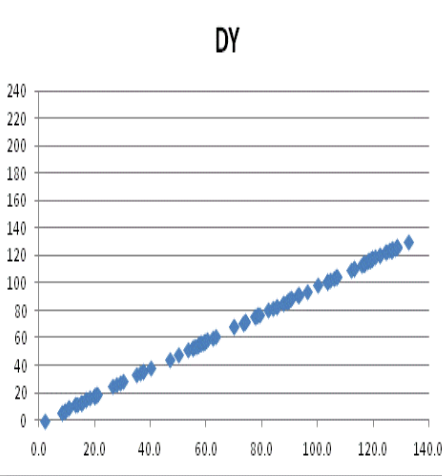
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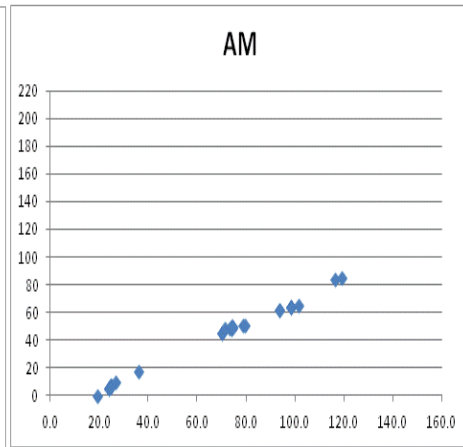
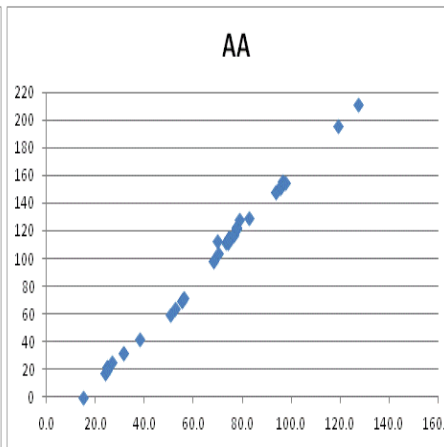
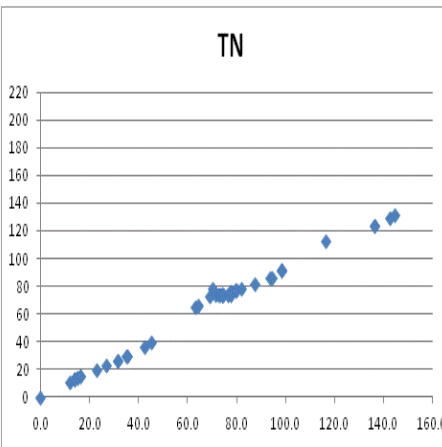
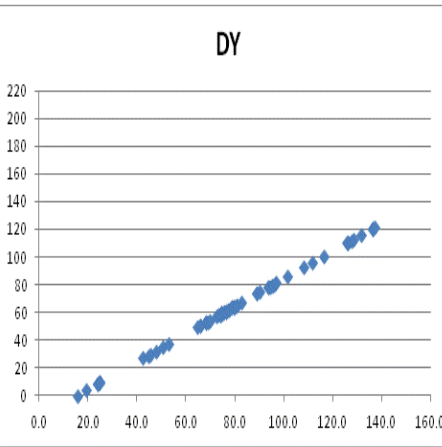
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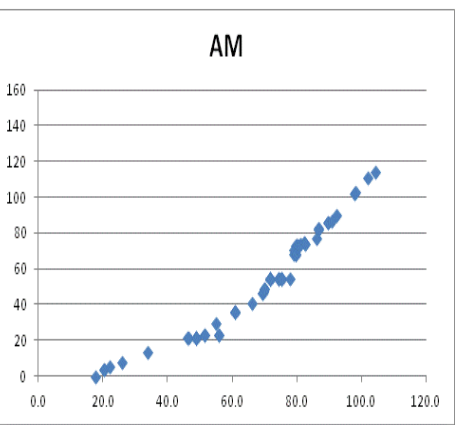
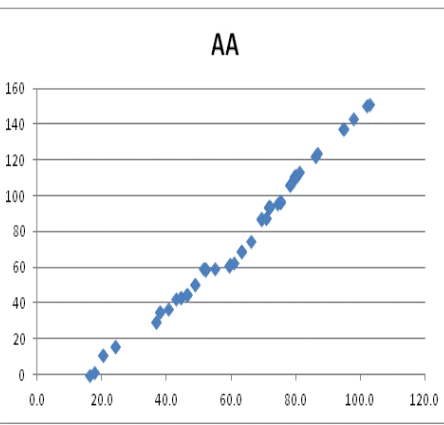
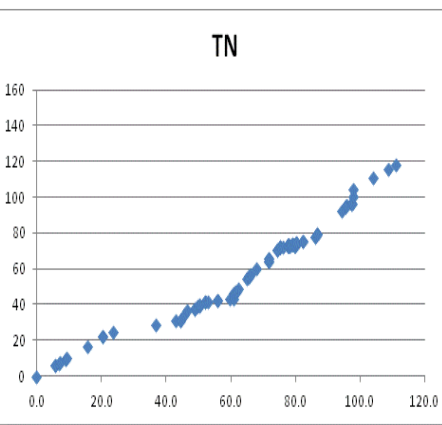
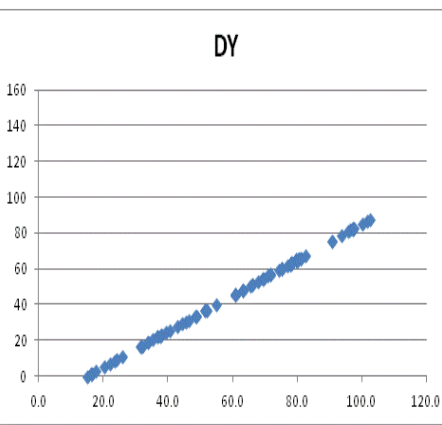
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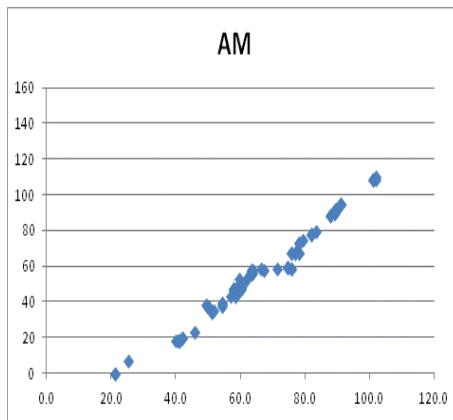
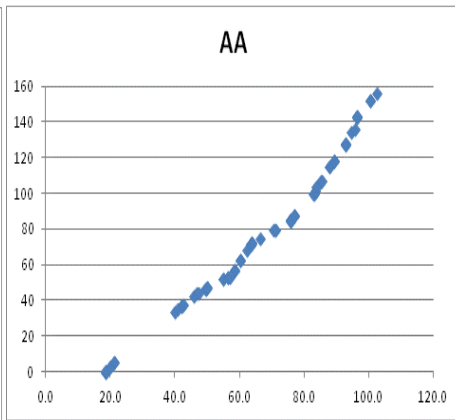
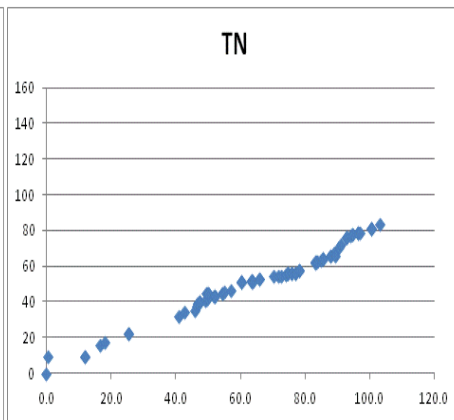
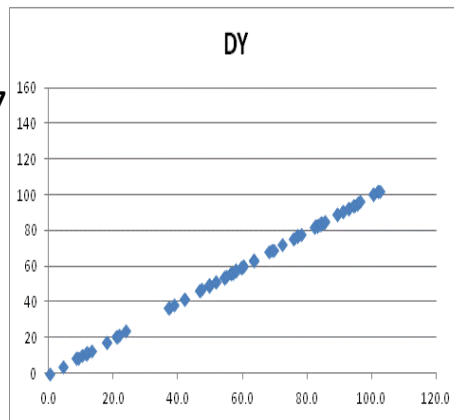
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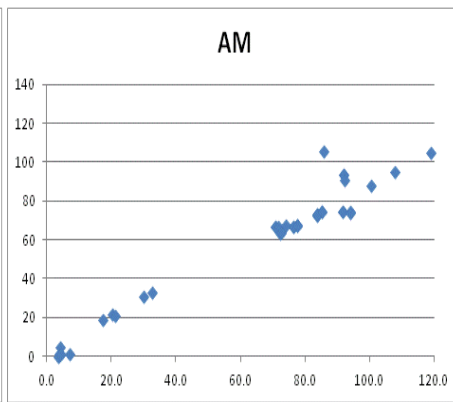
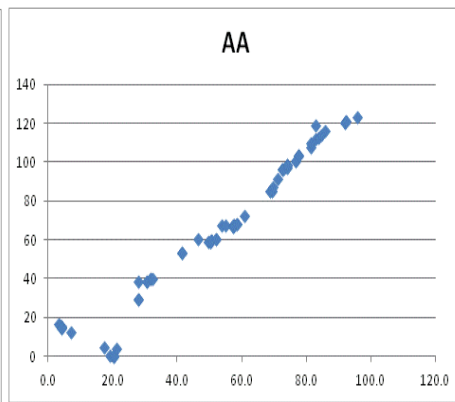
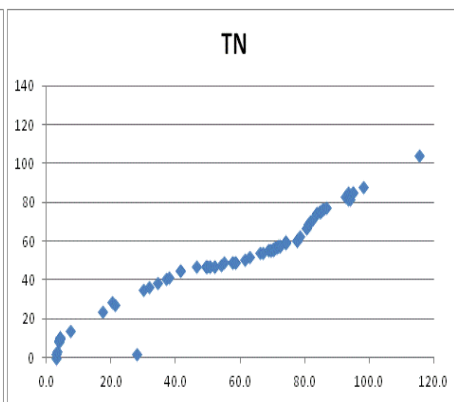
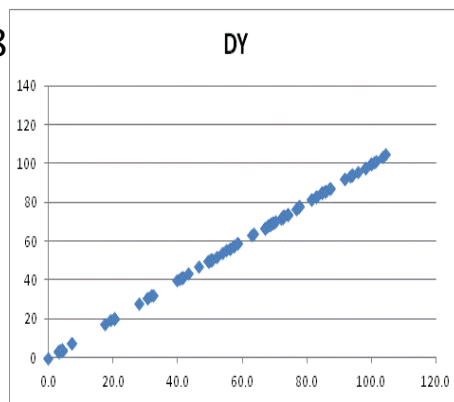
C6



C7



C8



C9

