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Linear Orbit Correction in the AGS to RHIC Transfer Lines

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Linear Orbit Correction in the AGS to RHIC Transfer Lines

T. Satogata

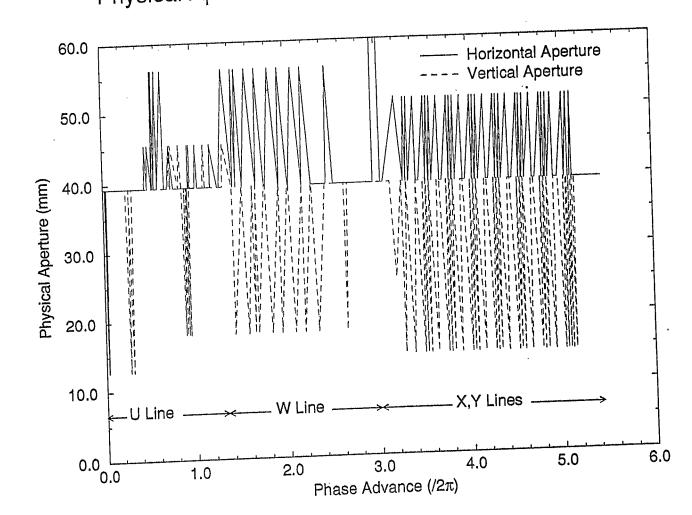
I. Introduction and Transfer Line Linear Correction System Layout

The AGS to RHIC transfer lines each cover a distance of approximately 600 meters, with phase advances of approximately 11 π radians in each plane. There are many vertical aperture restrictions present (mostly concentrated in the combined-function dipoles of the various bends), as well as a vertical pitch through the W line that makes these transfer lines non-planer. This investigation has several objectives, of which only one is reported on here. This is specifically to determine whether the linear orbit may be corrected to within one beam sigma in both planes, even with any single bpm or dipole corrector broken; more generally, where is the linear orbit correction in the AGS to RHIC transfer line best and where is it worst? The quantitative criterion may at first seem too stringent, but it must be stressed that adequate and robust linear orbit correction and optical matching in any transfer line are crucial for efficient beam transfer and flexibility of operation.

At the injection rigidity of $B\rho = 97.5 \text{ T} - \text{m}$ for beam extracted from the AGS, the sizes of gold and proton beams with typical emittances of 15π mm-mrad and 10π mm-mrad, respectively, are

$$\sigma(p) = 0.2833 \text{ mm} \sqrt{\beta[m]},
\sigma(Au) = 0.3639 \text{ mm} \sqrt{\beta[m]},$$
(1)

where β is the beta function at the observation point. β varies from about 4.5 m (U-line quad 10) to 175 m (U-line quad 9) in the horizontal plane, and from about 4 m (AGS extraction) to 100 m (U-line quad 4) in the vertical; in the X and Y 90 degree bends where vertical aperture is a serious consideration, the maximum betas are approximately 30 m. At these points the vertical aperture is 14.35 mm = $9.25 \sigma(p) = 7.20 \sigma(Au)$. The horizontal and vertical physical apertures are plotted in Figure (1), showing the vertical aperture restrictions of 14 to 17 mm in each of the major bends — eight degree, twenty degree and ninety degree. The horizontal aperture can be considered to be 39.37 mm throughout, with few constrictions; accurate horizontal steering is nonetheless important, especially at locations such as the switching magnet at the end of the W line and at injection into RHIC.



Physical Aperture of the AGS to RHIC Transfer Lines

Figure 1: Physical apertures of the AGS to RHIC transfer lines.

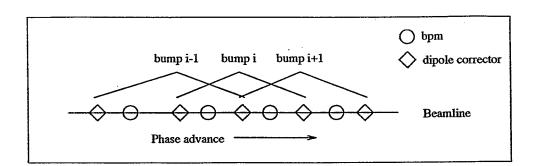


Figure 2: Overlapping 3-bumps in the clorb algorithm.

II. Clorb

Clorb is a general orbit correction routine which uses overlapping 3-bumps and a weighted penalty function to iteratively smooth the linear orbit. Because it does not rely on the specific form of the penalty function (which is traditionally RMS in other approaches), and because it can easily include nonlinear bpm and corrector response, including saturation and power supply limits, clorb is an extremely versatile correction algorithm. Originally used by Peggs et al. at CESR, it is also used in routine operation of the LBL ALS by Lindsay Schachinger and others. This section briefly describes this correction scheme as it will be applied in the AGS to RHIC transfer lines and RHIC.

Clorb is a linear orbit correction program that corrects the orbit based on a given set of position readings (typically BPMs), available dipole correctors and a linear orbit model (beta functions and phase advances). Clorb corrects each transverse plane independently, arranging correctors down the beamline it will correct in groups of three to create overlapping 3-bumps; this overlapping 3-bump layout is shown in Figure 2. For a transfer line this means that clorb corrects orbit displacements only between the first and last correctors it does not adjust the correctors at the end of the beamline to minimize downstream linear orbit errors, although it can be easily modified to do so. The penalty function of each bump is assessed for the three correctors and all bpms within the range of the bump. The relationship between bpm positions and corrector settings (or the bump strength) is calculated using either measured corrector-bpm response functions, as is done in the ALS, or a linear optics model of the beamline, as is done in this study.

The minimizing procedure of clorb is relatively simple: the penalty function is calculated for a bump and minimized by adjusting the bump strength. This is done in turn for each bump down the beamline and a total penalty function over all bumps is assessed; this procedure is then repeated, starting again with the first bump, until this global penalty function converges. This iterative approach to orbit correction, using a relatively simple model of the beamline, gives this scheme a robustness not found in other single-pass local-correction approaches. Clorb is not really a local correction algorithm (although it does correct locally) or a global correction algorithm (although it also minimizes a global penalty function) — it is in some sense both. Sets of bpms and correctors may be weighted differently for the penalty function to enforce tighter correction at critical locations; the penalty function may also return large penalties for corrector strengths that are out of range of their power supplies.

In the version of clorb used for AGS to RHIC transfer line orbit correction, a penalty function of the form

$$P = \sum_{i=1}^{N_{\text{bpms}}} B_i (X_i - X_{0,i})^2 + \sum_{j=1}^3 C_j , \qquad (2)$$

is used. B represents the bpm weight (1 for used bpms, 0 for broken or unused bpms), X and X_0 are the measured and intended beam positions (in mm), and C is the corrector penalty (1000 for an out of range corrector, 0 for an in range corrector). The units are arbitrary in this penalty function; a bump that needs no further correction has zero penalty. It was found during the course of this study that corrector strengths never saturated for realistic misalignment errors in the AGS to RHIC transfer line, and hence the second term in Equation (2) is negligible here.

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Care must be taken to avoid using three-bumps with anomalous distributions of bpms. Monitors that are closer than $\pi/10$ radians in betatron phase to correctors on either end of the three-bump are not used for penalty function calculation in that three-bump — it is presumed such a monitor will be corrected with other bumps. Monitors downstream of the last corrector that are not within any of the three-bumps used by clorb may also be corrected somewhat with the last pair of correctors in the beamline; this was not done in this study, and the deleterious effects of ignoring this correction can be seen in the results described in Section V.

Due to the nature of the design of the AGS to RHIC transfer line, it is important to turn off the corrector 90 degrees upstream of a monitor when it is known that monitor is broken; if this is not done the upstream corrector fight other correctors at unfavorably phased monitors, leading to unusually large corrector strengths and large anomalous bumps at locations without monitors. This approach is taken in Section V when the robustness of the linear orbit correction scheme is tested by turning off individual bpms.

III. Simulations without Randomly Generated Errors

Traditionally all magnetic and misalignment errors in a beamline are generated with random seeds, generating an ensemble of beamlines with "realistic" errors. After correction of each of these random lattices, statistics can be performed on the ensemble to determine the effectiveness of the correction scheme. Here we present an alternative scheme which may be used when the system under investigation is linear (for example, linear coupling and linear orbit correction) and the number of errors to be modeled and corrected is moderately small. This method gives exact results and does not use random seeds, thus eliminating drawbacks such as statistical scalings and sensitivities to anomalous seeds.

It is important to remember that the objective of either approach is to find the one-sigma linear orbit displacement σ^{orb} from the design trajectory, given a statistical description of one-sigma linear errors within the beamline. We shall show the equivalence of two approaches to this end. One option is to use an ensemble of N_{seeds} corrected beamlines, with randomly generated errors at all error locations for every beamline from seeds, and calculate the one-sigma linear orbit displacement σ_j^{orb} at any point j. If $x_{j,seed i}$ is the j^{th} position in the beamline characterized by seed i, then

$$\sigma_j^{orb} = \lim_{N_{seeds} \to \infty} \sqrt{\sum_{i=1}^{N_{seeds}} \frac{(x_{j, seed i} - \langle x \rangle_j)^2}{N_{seeds}}}, \qquad (3)$$

$$\langle x \rangle_j = \frac{1}{N_{seeds}} \sum_{i=1}^{N_{seeds}} x_{j,\text{seed i}}$$
 (4)

Alternatively, we can take an ensemble of N_{errs} corrected beamlines, where N_{errs} is the number of beamline locations with steering errors; each of these beamlines has a single one-sigma error at one of these locations and no steering errors elsewhere. The one-sigma linear orbit displacement is then calculated via

$$\sigma_j^{orb} = \sqrt{\sum_{i=1}^{N_{errs}} (x_{j,error\,i} - \langle x \rangle_j)^2} , \qquad (5)$$

$$\langle x \rangle_j = \frac{1}{N_{errs}} \sum_{i=1}^{N_{errs}} x_{j,\text{error i}}$$
 (6)

For errors that do not saturate dipole correctors during the correction procedure, it is reasonable to assume that the correction scheme used by clorb is linear with respect to superposition. That is, if a dipole error $C_{i,1}$ leads to an orbit displacement of $B_{j,1}$ after correction (and the same for a dipole error $C_{i,2}$ and corrected displacement $B_{j,2}$), then a dipole error $C_{i,1} + C_{i,2}$ will lead to an orbit displacement of $B_{j,1} + B_{j,2}$ after correction. With this assumption, the above formulae can be used to calculate the linear orbit distortion both before and after the correction procedure clorb has been applied.

The equivalence of Equations (5) and (6) to Equations (3) and (4) can be seen from the following reasoning. Because the orbit correction scheme is local, the sums in (5) and (6) for any point x_j in the beamline only carry local contributions; any errors located far upstream or downstream of this point contribute little. This cancellation is performed in (3) and (4) by a phase averaging of many ensembles of errors, all approximately cancelling one another. Since this system is linear with respect to superposition, the averaging performed by taking random beamlines can instead be performed explicitly — we know that errors far away will *exactly* cancel, and the only errors left contributing are local errors. The effect of these local errors can then be summed independently (again assuming superposition) to arrive at Equations (5) and (6).

Therefore, instead of taking the RMS values of the linear orbit over an ensemble of randomly generated (corrected) beamlines, we can take the root of the sum of the squares of 1-sigma orbits (again corrected) over the ensemble of all single 1-sigma errors. There is no scaling by the square root of the number of seeds in the latter, and this method is completely deterministic. These methods are compared in the Section V, where they are found to be in close agreement.

IV. Simulation Procedure and Results

The simulation procedure is diagramed in Figure (3), where processes are represented by lozenges and data are represented by rectangles. Arrows indicate directions of data flow. The machine simulation comprises the left-hand side of the figure, where random dipole errors are simulated; their effects on the orbit are calculated in lorb, a linear orbit simulation module. Lorb gives a simulated linear orbit at *all* elements in the beamline ("God's linear orbit") which may be displayed, saved and analyzed.

The central column represents information in the optics database, including information about the lattice layout and elements (in the Lattice SDS) and the beta functions and phase advances between various elements in the lattice (in the Twiss SDS). The linear lattice used in this study is the design lattice, as described in the "Holy Lattice" area used by the RHIC Accelerator Physics group. It is important to note that the optical parameters used are design values, and there is no effort to feed errors or misalignments

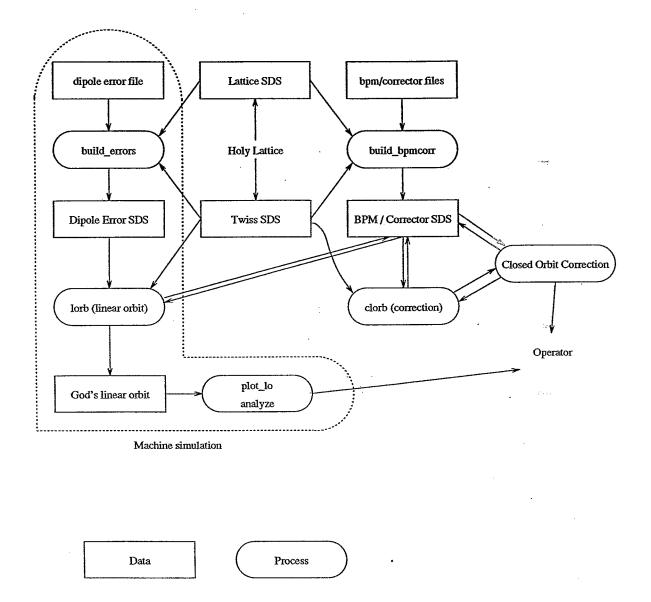


Figure 3: Simulation procedure for AGS to RHIC Transfer Line study, with arrows representing the directions of data flow.

back into this database. Thus this study uses a base design model of the machine plus small well-controlled perturbations from that design.

The left column represents specific information about BPMs and dipole correctors that are used for the orbit correction process. A list of BPMs and correctors to use is retrieved from a file, and data structures are set up in the form of an SDS in shared memory. The simulation of orbit correction is then performed:

• Initialize the BPM and dipole corrector data structures.

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- Read the set of dipole errors, either randomly generated from a seed at all misalignment points in the lattice, or a single one-sigma magnet misalignment error as described in the previous section.
- Generate a distorted linear orbit at all BPM locations with lorb.
- Use clorb to correct this distorted orbit at all BPM locations, using the listed dipole correctors. Typically several runs of clorb are necessary here for convergence.
- Use lorb to generate a corrected linear orbit at all magnet positions. Store this linear orbit for later analysis.

This procedure allows the simulation of many types of steering error and their correction. There are a variety of sources of these errors, including quadrupole misalignments, dipole rolls, orbit errors in higher order multipoles and differences of actual and design magnet field strengths. In this paper we consider only the first two sources. Nick Tsoupas has recently investigated the last source in the 90 degree bends, based on field variations between new and old steel dipole magnets reported by Peter Wanderer, and found corrector tolerances in this situation to be quite acceptable.

A quadrupole of strength k_1 and length L gives a transverse kick of $\Delta x' = -k_1 L x$; a transverse misalignment of this quad by \bar{x} then gives a steering error of

$$\Delta x'(\text{quad}) = -k_1 L \,\bar{x} \tag{7}$$

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in either dimension. Similarly, the vertical kick from a dipole with bend angle θ rolled by ψ_{roll} around the horizontal axis is approximately

$$\Delta x'(\text{dipole}) = \theta(1 - \cos \psi_{roll})$$

$$\Delta y'(\text{dipole}) = \theta \sin \psi_{roll}.$$

This is not strictly correct, but serves as an adequate approximation for this study. Here each dipole was assumed to have an RMS survey roll error of 1.0 mrad, and each quadrupole had an RMS survey transverse displacement of 0.5 mm; these values are somewhat higher than the tolerances expected from survey in these transfer lines.

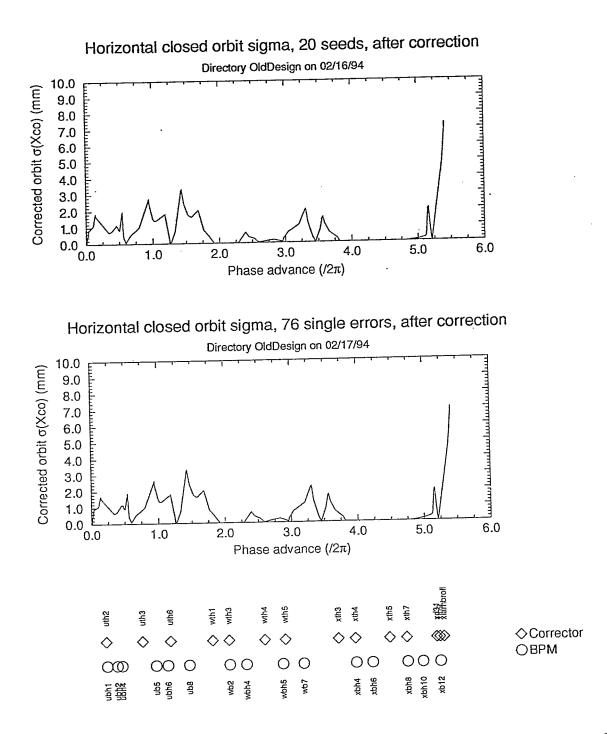
V. Results and Conclusions

Several tests of correction robustness were applied to the AGS to RHIC transfer lines. First, the misaligned transfer line was simulated using either 20 seeds or 76 single deterministic errors, and the resulting one-sigma orbit displacements were compared. These results are shown in Figures (4) and (5), and there is little noticeable difference between the two methods. Each of these figures also shows the corrector and BPM layout for each plane in the transfer lines (notable is the two-corrector two-BPM pairing design), as well as placement of flags which can be used as surrogate BPMs if necessary for beamline orbit tuning. The similarity of these corrected linear orbits demonstrates that the reasoning in Section III is correct, and an analysis of the beamline orbit correction can proceed without any further reference to seeds, randomly generated errors and statistical weighting.

It is also important to note that in these figures the orbit correction appears to be faulty at a crucial point in the beamline, where the beam goes through the injection lambertson and into RHIC. The lack of correction by clorb in this area was mentioned in Section II, and in the future clorb will be altered to adjust correctors at the end of the transfer line and steer beam properly through the lambertson and injection region.

To test the robustness of the linear orbit correction system against breakdown, single BPMs were broken and the correction then applied to see where additional system redundancy might be required. There are several locations in the transfer lines which might suffer greatly if a single BPM plane were to break — the wbpm2/wbpm4 pair, the xbpmh4/xbpmh5 pair, and xbpmv3 to name a few. The AGS to RHIC transfer line linear orbit correction system was designed so that the orbit could be completely corrected at several points through the line by pairs of correctors and BPMs placed 90 degrees out of phase.

For completeness, single correctors were also broken to test the correction system. In these circumstances clorb corrected the linear orbit of the beamline nearly as well as with all correctors working, demonstrating that the design of the linear orbit correction system is quite robust when single correctors are broken. There were no correctors in either plane



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Figure 4: A comparison of horizontal linear orbit correction in the AGS to RHIC transfer line with and without random error seeds. Steering errors are quadrupole misalignments (0.5 mm RMS) and dipole rolls (1.0 mrad RMS).

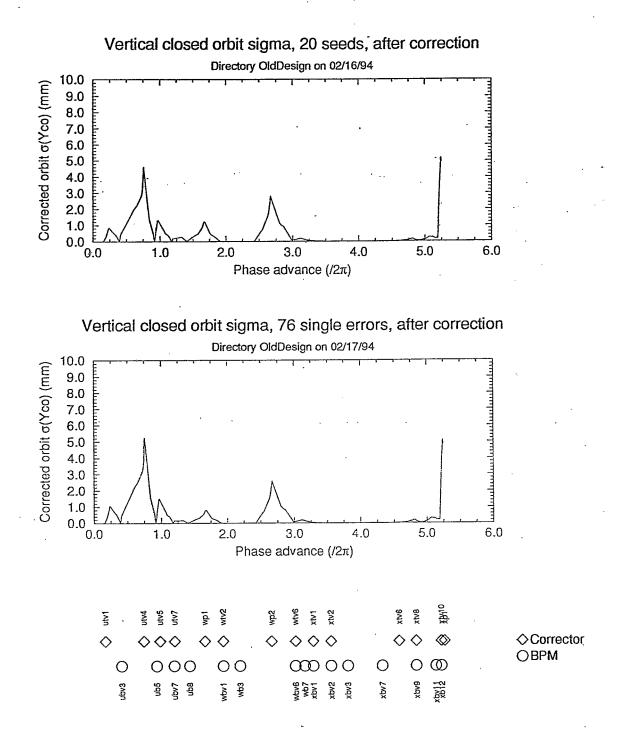


Figure 5: Same as previous figure, for vertical linear orbit correction.

that seemed critical or that led to an unusual inability of the correction system to correct a distorted linear orbit.

In the horizontal plane one critical pair of BPMs are wbpmh2 and wbpmh4, located at horizontal phase advances of approximately 2.00 and 2.25 down the beamline. These BPMs are crucial for correction of upstream steering errors upon entering the W line, and their placement near the end of the 20 degree bend also requires their presenence to diagnose any steering problems in this arc. Figure (6) shows that the horizontal orbit errors accumulated over the previous 270 degrees in phase cannot be fully corrected with either one of these BPMs broken, leading to one sigma RMS orbit errors after correction of 10-20 mm. There is no flag nearby for use as a backup position monitor. This area is also of some concern in vertical steering, since the W line contains a vertical drop between pitching magnets wp1 and wp2. Failure of the wbpmv1 BPM plane leads to corrected one sigma RMS orbit errors of up to 15 mm just downstream (Figure 7), within the dipole magnets of the 20 degree bend, so there is also a need for another vertical position monitor in this area.

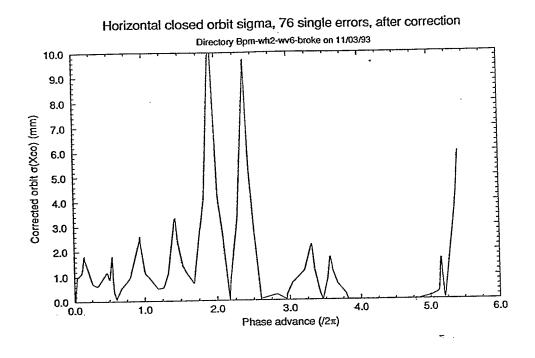
Another crucial area for orbit correction in both transverse planes is the entry into the X and Y lines, or 90 degree bends. Examining Figures (4) and (5), we see several interesting features. In the horizontal plane, pairs of BPMs are isolated (xbpmh4 and xbpmh5, and xbpmh7 and xbpmh9) without backup readings available. In the vertical plane where the aperture restriction is severe, there is a long region between xbpmv1 and xbpmv8 where several BPMs are placed approximately 180 degrees apart — if xbpmv2 were to break, there would be no information about the phase of the linear orbit that it covered until the end of the arc. The effects on the corrected orbit from each of these BPMs breaking can be seen in Figures (8) through (10); of serious interest is the orbit oscillation in Figure (10) when BPM xbpmv2 is broken, showing that phase coverage of the orbit correction system is insufficient at that point.

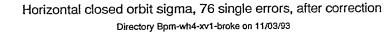
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For adequate phase coverage at these critical points (through the 20 and 90 degree bends), a rather minimal number of BPM planes can be added. The BPMs wbpmh2 and wbpmh3, previously single-plane, can each be converted to dual-plane BPMs. In this sense each provides phase-redundancy for both BPMs in each plane that bracket them, and allow for full orbit correction even in the event of failure of any single BPM. A dual-plane BPM can also be added in each of the X and Y lines between dipoles D12 and D13. This BPM provides additional horizontal coverage (especially useful considering that horizontal correction may be necessary to compensate for old- and new-steel gradient dipole differences) as well as the phase coverage necessary for redundant correction of the vertical orbit. This

location does not contain any instrumentation or restrictions that would interfere with the installation of an additional dual-plane BPM.

Following is a memo detailing these changes, later implemented by Ted Robinson and Waldo MacKay.





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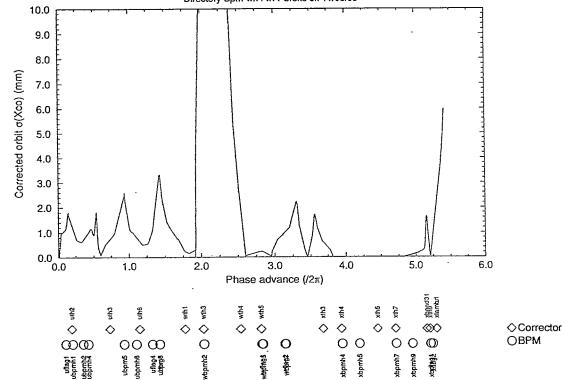


Figure 6: The corrected horizontal linear orbit with BPMs wbpmh2 and wbpmh4 each individually broken.

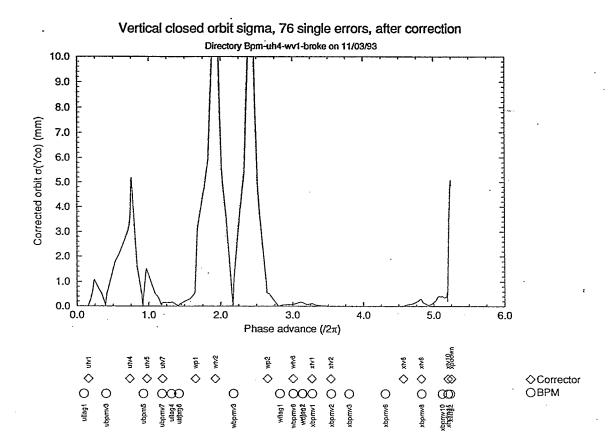


Figure 7: The corrected vertical linear orbit with BPM wbpmv1 broken.

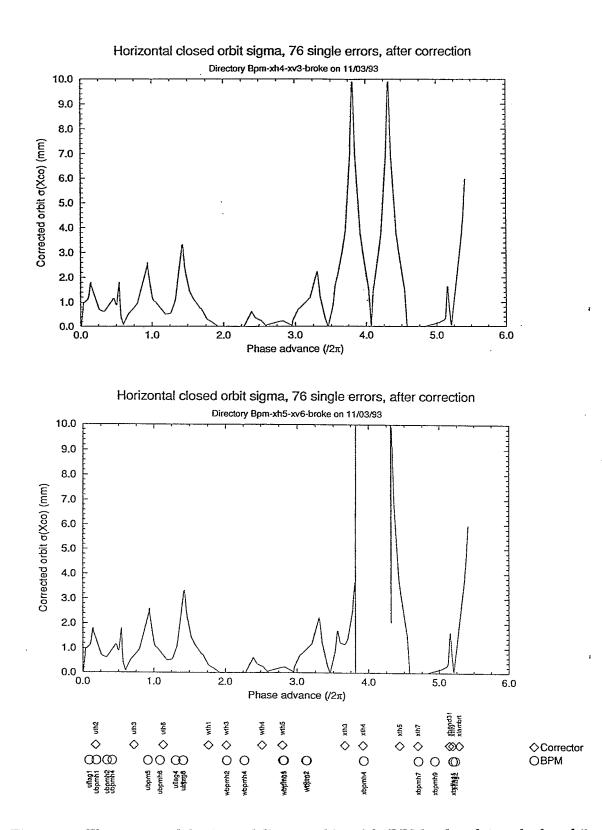
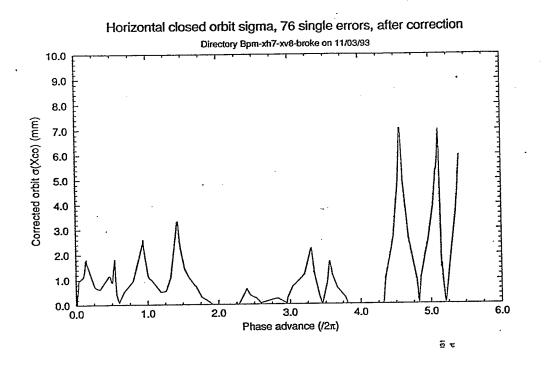


Figure 8: The corrected horizontal linear orbit with BPMs xbpmh4 and xbpmh5 each individually broken.



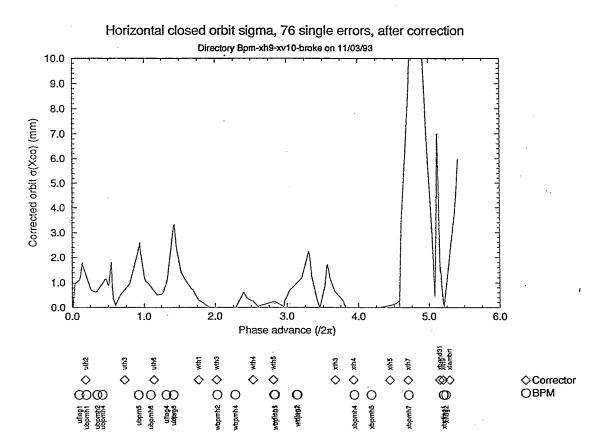
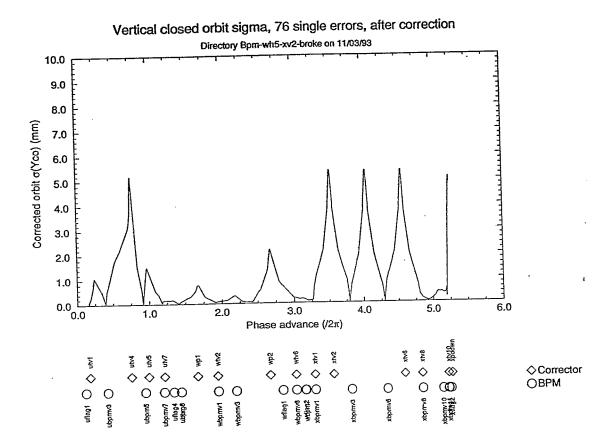


Figure 9: The corrected horizontal linear orbit with BPMs xbpmh7 and xbpmh9 each individually broken.



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Figure 10: The corrected vertical linear orbit with BPM xbpmv2 broken, showing the orbit oscillation from lack of sufficient phase coverage.

Appendix: Memo detailing BPM layout changes

Changes in BPM layout in the AGS-RHIC Transfer Lines 11/12/93

T. Satogata

To: H. Foelsche, T. Robinson, T. Shea CC: W. MacKay, S. Peggs, P. Zhou

The current organization of BPMs and dipole correctors in the AGS to RHIC transfer lines has been studied using a linear correction model that will be implemented during the sextant test. A set of typical quadrupole misalignments (feeding down to dipole errors) and dipole roll errors have been used to predict the one-sigma closed orbit after linear orbit correction. This memo details the resulting changes in BPM layout through these transfer lines; there are no changes in the dipole corrector layout, as the current configuration of these elements is adequate.

The criterion for adequate correction of the linear orbit is that, given typical errors, the aperture at all points in the transfer line should be at least 3 sigma of the closed orbit error after correction. This criterion should be met even when any single BPM is broken for adequate redundancy of the correction system. Redundancy of information is also necessary for adequate commissioning of the beamline and controls system, where discrepancies are used to discover errors in installation, and weaknesses and strengths of the design.

The entry into the W line contains a pair of dipole correctors and BPMs in each plane, placed 90 degrees apart in phase advance and designed to fully correct the beam orbit as it enters this region. These pairs are:

Horizontal Correctors: wth1, wth3 Horizontal BPMs: wbpmh2, wbpmh4

Vertical Correctors: wp1, wtv2 Vertical BPMs: wbpmv1, wbpmv3

Failure of any one of these BPMs greatly reduces the effectiveness of orbit correction at this point, in either plane. No flags are nearby for use as backup BPMs, but BPMs wbpmh2 and wbpmv3 may be converted to dual-plane BPMs with acceptable phase coverage. Note that this conversion requires no additional vacuum penetrations or change in BPM numbering.

The major aperture restriction of the entire transfer line is the 1.4 cm (radius) vertical aperture in the 90 degree bends (X and Y lines). This following discussion applies to both these bends. The current design for vertical BPM placement in and around the upstream end of these bends has three BPMs placed approximately 180 degrees apart: xbpmv1, xbpmv3, and xbpmv6. These BPMs are redundant, in that each gives essentially the same phase information about the linear orbit. The critical BPM is xbpmv2, which has no redundant BPM in phase with it until the end of the transfer line. This is unacceptable failure of this BPM results in a 1-sigma closed orbit error, after correction, of 6-7 mm through the bend.

There is also a lack of redundancy in the horizontal plane in the same region of the 90 degree bends; BPMs xbpmh4 and xbpmh5 alone provide orbit information. Both this and the vertical BPM problem are resolved by placing an additional dual-plane BPM in the straight section between dipoles D12 and D13, in both the X and Y lines. At present this straight contains only an ion pump, which will not interfere with the performance of a nearby BPM and which leaves sufficient space for such an installation.

Table 1: BPM placement in AGS-RHIC Transfer Line As of 11/12/93

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Name	plane	s-loc (end)	mu_x	mu_y
ubpmh1	· h	5.149	0.014	0.188
ubpmh2	h	16.825	0.040	0.339
ubpmv3	v	33.325	0.119	0.395
ubpmh4	h	37.983	0.147	0.412
ubpm5	hv	88.016	0.566	0.918
ubpmh6	h	112.601	0.938	1.089
ubpmv7	v	124.090	1.005	1.179
ubpm8	hv	159.003	1.262	1.412
wbpmv1	v	230.642	1.773	1.920
wbpmh2	h	248.351	1.922	2.021
wbpmv3	v	266.061	2.022	2.172
wbpmh4	h	283.771	2.172	2.274
- wbpmh5	h	348.186	2.601	2.822
wbpmv6	v	368.507	2.801	2.998
wbpm7	hv	389.400	2.955	3.134
xbpmv1	v	403.095	3.056	3.265
xbpmv2	v	425.171	3.447	3.534
- xbpmv3	v	443.313	3.688	3.787
xbpmh4	h	453.089	3.817	3.928
xbpmh5	h	471.231	4.067	4.172
xbpmv6	v	480.537	4.194	4.300
xbpmh7	h	508.925	4.577	4.697
- xbpmv8	v	518.231	4.713	4.815
- xbpmh9	h	526.683	4.825	4.930
xbpmv1	0 v	543.459	5.085	5.112
xbpm11	hv	561.084	5.211	5.196

ybpmv1	v	403.095	3.056	3.265
ybpmv2	v	425.171	3.447	3.534
ybpmv3	v	443.313	3.688	3.787
ybpmh4	h	453.089	3.817	3.928
ybpmh5	h	471.231	4.067	4.172
ybpmv6	v	480.537	4.194	4.300
ybpmh7	h	508.925	4.577	4.697
ybpmv8	v	518.231	4.713	4.815 ·
ybpmh9	h	526.683	4.825	4.930
ybpmv10	v	543.459	5.085	5.112
ybpm11	hv	561.084	5.211	5.196

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Nameplanes-loc (end) mu_x mu_y ubpmh1h5.1490.0140.188ubpmh2h16.8250.0400.339ubpmv3v33.3250.1190.395ubpmh4h37.9830.1470.412ubpmb5hv88.0160.5660.918ubpmk4h112.6010.9381.089ubpmk7v124.0901.0051.179ubpm8hv159.0031.2621.412wbpm1v230.6421.7731.920wbpm3hv266.0612.0222.172 < change from H to dualwbpm44h283.7712.1722.274wbpmb5h348.1862.6012.822wbpm7hv389.4002.9553.134xbpmv1v403.0953.0563.265xbpmv1v403.0953.0563.265xbpmv1v403.0953.0563.265xbpmv3v443.3133.6883.787xbpm5hv462.1553.9804.058 < added dual planexbpm5hv462.1553.9804.058 < added dual planexbpm5h368.9254.6774.697xbpm4h453.0374.1944.300xbpm5hv462.1553.9804.058 < added dual planexbpm5hv462.1553.9804.057xbpm6h471.2314.0674.172		itions a	acement in AGS-R nd changes	HIC Transf	er Line
ubpmh2 h 16.825 0.040 0.339 ubpmv3 v 33.325 0.119 0.395 ubpmk4 h 37.983 0.147 0.412 ubpm5 hv 88.016 0.566 0.918 ubpm6 h 112.601 0.938 1.089 ubpmv7 v 124.090 1.005 1.179 ubpmv1 v 124.090 1.005 1.179 ubpmv1 v 230.642 1.773 1.920 wbpm2 hv 248.351 1.922 2.021 < change from H to dual wbpm3 hv 266.061 2.022 2.172 < change from V to dual wbpm4 h 283.771 2.172 2.274 wbpm45 h 348.186 2.601 2.822 wbpm47 hv 368.507 2.801 2.998 wbpm47 hv 403.095 3.056 3.265 xbpmv4 h 453.089 3.817 3.928	Name	plane	s-loc (end)	mu_x	 mu_y
ubpmv3v33.3250.1190.395ubpmh4h37.9830.1470.412ubpm5hv88.0160.5660.918ubpm6h112.6010.9381.089ubpmv7v124.0901.0051.179ubpm8hv159.0031.2621.412vv230.6421.7731.920vbpm2hv248.3511.9222.021 < change from H to dual	ubpmh1	h	5.149	0.014	0.188
ubpmh4h37.9830.1470.412ubpm5hv88.0160.5660.918ubpm6h112.6010.9381.089ubpm7v124.0901.0051.179ubpm8hv159.0031.2621.412vv230.6421.7731.920wbpm2hv248.3511.9222.021 < change from H to dual	ubpmh2	h	16.825	0.040	0.339
ubpm5hv88.0160.5660.918ubpmh6h112.6010.9381.089ubpm7v124.0901.0051.179ubpm8hv159.0031.2621.412wbpm1v230.6421.7731.920wbpm2hv248.3511.9222.021 < change from H to dual	ubpmv3	v	33.325	0.119	0.395
ubpmh6h112.6010.9381.089ubpm7v124.0901.0051.179ubpm8hv159.0031.2621.412vbpm1v230.6421.7731.920wbpm2hv248.3511.9222.021 < change from H to dual	ubpmh4	h	37.983	0.147	0.412
ubpmv7v124.0901.0051.179ubpm8hv159.0031.2621.412wbpmv1v230.6421.7731.920wbpm2hv248.3511.9222.021 < change from H to dual	ubpm5	hv	88.016	0.566	0.918
ubpm8hv159.0031.2621.412vbpmv1v230.6421.7731.920wbpm2hv248.3511.9222.021 < change from H to dual	ubpmh6	h	112.601	0.938	1.089
wbpmv1v230.6421.7731.920wbpm2hv248.3511.9222.021 < change from H to dualwbpm3hv266.0612.0222.172 < change from V to dualwbpm4h283.7712.1722.274wbpmb5h348.1862.6012.822wbpmv6v368.5072.8012.998wbpm7hv389.4002.9553.134xv403.0953.0563.265xbpmv1v403.0953.6883.787xbpmv3v443.3133.6883.787xbpm4h453.0893.8173.928xbpm5hv462.1553.9804.058 < added dual planexbpmk6h471.2314.0674.172xbpmN8h508.9254.5774.697xbpm9v518.2314.7134.815	ubpmv7	v .	124.090	1.005	1.179
wbpm2hv248.3511.9222.021 < change from H to dualwbpm3hv266.0612.0222.172 < change from V to dualwbpmh4h283.7712.1722.274wbpmb5h348.1862.6012.822wbpmv6v368.5072.8012.998wbpm7hv389.4002.9553.134v403.0953.0563.265xbpmv1v403.0953.6883.787xbpmv3v443.3133.6883.787xbpm4h453.0893.8173.928xbpm5hv462.1553.9804.058 < added dual planexbpmh6h471.2314.0674.172xbpmt7v480.5374.1944.300xbpm8h508.9254.5774.697xbpm9v518.2314.7134.815	ubpm8	hv	159.003	1.262	1.412
wbpm2hv248.3511.9222.021 < change from H to dualwbpm3hv266.0612.0222.172 < change from V to dualwbpmh4h283.7712.1722.274wbpmb5h348.1862.6012.822wbpmv6v368.5072.8012.998wbpm7hv389.4002.9553.134v403.0953.0563.265xbpmv1v403.0953.6883.787xbpmv3v443.3133.6883.787xbpm4h453.0893.8173.928xbpm5hv462.1553.9804.058 < added dual planexbpmh6h471.2314.0674.172xbpmt7v480.5374.1944.300xbpm8h508.9254.5774.697xbpm9v518.2314.7134.815					
wbpm3 hv 266.061 2.022 2.172 < change from V to dual	wbpmv1	v	230.642	1.773	1.920
wbpmh4h283.7712.1722.274wbpmh5h348.1862.6012.822wbpmv6v368.5072.8012.998wbpm7hv389.4002.9553.134xbpmv1v403.0953.0563.265xbpmv2v425.1713.4473.534xbpmv3v443.3133.6883.787xbpm4h453.0893.8173.928xbpm5hv462.1553.9804.058 < added dual planexbpmh6h471.2314.0674.172xbpmk7v480.5374.1944.300xbpmh8h508.9254.5774.697xbpmv9v518.2314.7134.815	wbpm2	hv	248.351	1.922	2.021 < change from H to dual
wbpmh5h348.1862.6012.822wbpmv6v368.5072.8012.998wbpm7hv389.4002.9553.134xbpmv1v403.0953.0563.265xbpmv2v425.1713.4473.534xbpmv3v443.3133.6883.787xbpm4h453.0893.8173.928xbpm5hv462.1553.9804.058 < added dual planexbpm6h471.2314.0674.172xbpm8h508.9254.5774.697xbpm49v518.2314.7134.815	wbpm3	hv	266.061	2.022	2.172 < change from V to dual
wbpmv6v368.5072.8012.998wbpm7hv389.4002.9553.134xbpmv1v403.0953.0563.265xbpmv2v425.1713.4473.534xbpmv3v443.3133.6883.787xbpm4h453.0893.8173.928xbpm5hv462.1553.9804.058 < added dual planexbpmh6h471.2314.0674.172xbpmk7v480.5374.1944.300xbpm8h508.9254.5774.697xbpmv9v518.2314.7134.815	wbpmh4	h. h	283.771	2.172	2.274
wbpm7hv389.4002.9553.134xbpmv1v403.0953.0563.265xbpmv2v425.1713.4473.534xbpmv3v443.3133.6883.787xbpm4h453.0893.8173.928xbpm5hv462.1553.9804.058 < added dual plane	wbpmh5	h	348.186	2.601	2.822
xbpmv1v403.0953.0563.265xbpmv2v425.1713.4473.534xbpmv3v443.3133.6883.787xbpmh4h453.0893.8173.928xbpm5hv462.1553.9804.058 < added dual planexbpmh6h471.2314.0674.172xbpmv7v480.5374.1944.300xbpmh8h508.9254.5774.697xbpmv9v518.2314.7134.815	wbpmv6	v	368.507	2.801	2.998
xbpmv2v425.1713.4473.534xbpmv3v443.3133.6883.787xbpmh4h453.0893.8173.928xbpm5hv462.1553.9804.058 < added dual planexbpmh6h471.2314.0674.172xbpmv7v480.5374.1944.300xbpmh8h508.9254.5774.697xbpmv9v518.2314.7134.815	wbpm7	hv	389.400	2.955	3.134
xbpmv2v425.1713.4473.534xbpmv3v443.3133.6883.787xbpmh4h453.0893.8173.928xbpm5hv462.1553.9804.058 < added dual planexbpmh6h471.2314.0674.172xbpmv7v480.5374.1944.300xbpmh8h508.9254.5774.697xbpmv9v518.2314.7134.815					
xbpmv3v443.3133.6883.787xbpmh4h453.0893.8173.928xbpm5hv462.1553.9804.058 < added dual planexbpmh6h471.2314.0674.172xbpmv7v480.5374.1944.300xbpmh8h508.9254.5774.697xbpmv9v518.2314.7134.815	xbpmv1	v	403.095	3.056	3.265
xbpmh4h453.0893.8173.928xbpm5hv462.1553.9804.058 < added dual planexbpmh6h471.2314.0674.172xbpmv7v480.5374.1944.300xbpmh8h508.9254.5774.697xbpmv9v518.2314.7134.815	xbpmv2	v	425.171	3.447	3.534
xbpm5hv462.1553.9804.058 < added dual planexbpmh6h471.2314.0674.172xbpmv7v480.5374.1944.300xbpmh8h508.9254.5774.697xbpmv9v518.2314.7134.815	xbpmv3	v	443.313	3.688	3.787
xbpmh6 h 471.231 4.067 4.172 xbpmv7 v 480.537 4.194 4.300 xbpmh8 h 508.925 4.577 4.697 xbpmv9 v 518.231 4.713 4.815	xbpmh4	h	453.089	3.817	3.928
xbpmv7v480.5374.1944.300xbpmh8h508.9254.5774.697xbpmv9v518.2314.7134.815	xbpm5	hv	462.155	3.980	4.058 < added dual plane
xbpmh8h508.9254.5774.697xbpmv9v518.2314.7134.815	xbpmh6	h	471.231	4.067	4.172
xbpmv9 v 518.231 4.713 4.815	xbpmv7	v	480.537	4.194	4.300
-	xbpmh8	h.	508.925	4.577	4.697
xbpmh10 h 526.683 4.825 4.930	xbpmv9	v	518.231	4.713	4.815
	xbpmh10) h	526.683	4.825	4.930

xbpmv11 xbpm12	v hv	543.459 561.084	5.085 5.211	5.112 5.196
ybpmv1	v	403.095	3.056	3.265
ybpmv2	v	425.171	3.447	3.534
ybpmv3	v	443.313	3.688	3.787
ybpmh4	h	453.089	3.817	3.928
ybpm5	hv	462.155	3.980	4.058 < added dual plane
ybpmh6	h	471.231	4.067	4.172
ybpmv7	v	480.537	4.194	4.300
ybpmh8	h	508.925	4.577	4.697
ybpmv9	v	518.231	4.713	4.815
ybpmh10	h	526.683	4.825	4.930
ybpmv11	v	543.459	5.085	5.112
ybpm12	hv	561.084	5.211	5.196

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